

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
31 October 2002 (31.10.2002)

PCT

(10) International Publication Number
WO 02/085933 A1

(51) International Patent Classification⁷: **C07K 14/255**,
C12N 15/31, A61K 39/00

D. [US/US]; 14038 23rd Avenue N.E., Seattle, WA 98125 (US).

(21) International Application Number: PCT/US01/22978

(74) Agents: **GAY, David, A. et al.**; Campbell & Flores LLP, 4370 La Jolla Village Drive, San Diego, CA 92122 (US).

(22) International Filing Date: 20 July 2001 (20.07.2001)

(81) Designated States (*national*): AE, AG, AL, AM, AT (utility model), AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ (utility model), CZ, DE (utility model), DE, DK (utility model), DK, DM, DZ, EC, EE (utility model), EE, ES, FI (utility model), FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK (utility model), SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/285,477 20 April 2001 (20.04.2001) US

(63) Related by continuation (CON) or continuation-in-part (CIP) to earlier application:
US 60/285,477 (CON)
Filed on 20 April 2001 (20.04.2001)

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

(71) Applicant (*for all designated States except US*): **THE INSTITUTE FOR SYSTEMS BIOLOGY** [US/US]; 4225 Roosevelt Way, N.E., Suite 200, Seattle, WA 98105-6099 (US).

Published:

- with international search report
- before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments

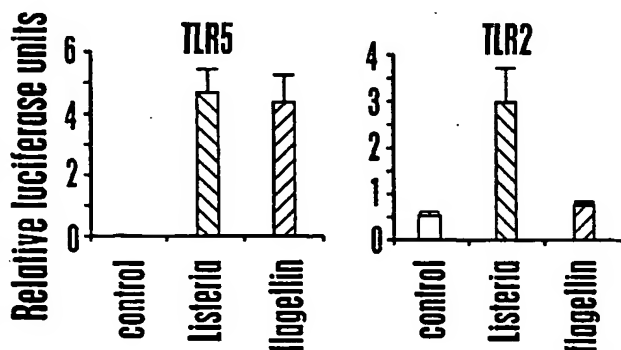
(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **ADEREM, Alan** [ZA/US]; 715 37th Avenue, Seattle, WA 98122 (US). **HAYASHI, Fumitaka** [US/US]; 95 West Squantum Street #405, North Quincy, MA 02171 (US). **UNDERHILL, David, M.** [US/US]; 4218 N.E. 89th Street, Seattle, WA 98115 (US). **OZINSKY, Adrian** [ZA/US]; 4332 Bagley, Avenue North, Seattle, WA 98103 (US). **SMITH, Kelly,**

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: TOLL-LIKE RECEPTOR 5 LIGANDS AND METHODS OF USE

WO 02/085933 A1



(57) Abstract: The invention provides an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence GAVQNRNFNSAIT, or a modification thereof, and having toll-like receptor 5 (TLR5) binding. Methods of inducing an immune response are also provided.

TOLL-LIKE RECEPTOR 5 LIGANDS AND METHODS OF USE

BACKGROUND OF THE INVENTION

Cancer is the second leading cause of death in the United States, accounting for one in every four
5 deaths. This year, it is expected that over 1500 Americans will die of cancer each day and that a million new cases of cancer will be diagnosed. The most common treatments for cancer are surgery, radiation and chemotherapy. According to the American Cancer Society,
10 immunotherapy can be considered as the "fourth modality" in the treatment of cancer. Immunotherapy is treatment that stimulates one's own immune system to fight cancer.

Cancer is a group of diseases characterized by uncontrolled growth of abnormal cells of the body. All
15 types of cancer involve the malfunction of genes that control cell growth and division. Some of these genes become incorrectly regulated, resulting in over- or under-production of a particular protein, while others become mutated, resulting in unusual or abnormal proteins
20 that alter normal cellular functions. These abnormal proteins, referred to as "tumor cell antigens," should be recognized and destroyed by an individual's immune system as "foreign" antigens.

However, the immune system of a cancer patient
25 may ignore these tumor antigens and be unresponsive to the growing tumor. Using immunotherapy approaches, such as cancer vaccines and immune system modulators, an individual's immune system can be induced to mount a potent immune response against tumor cell antigens,

resulting in elimination of cancer cells. A cancer vaccine can contain a tumor cell antigen that stimulates the immune system to recognize and destroy cells which display that antigen. Treating an individual with such a
5 cancer vaccine can result in a humoral response, which involves producing antibodies that recognize and target tumor cells for destruction and a cellular response, which involves producing cytotoxic T cells that recognize and destroy tumor cells directly, or both responses. It
10 can be desirable to obtain both a humoral and cellular immunity response during immunotherapy because both arms of immune response have been positively correlated with beneficial clinical responses. To help stimulate either or both humoral and cellular immune responses, a cancer
15 vaccine can be combined with an adjuvant, which is a substance that stimulates a general immune response.

The potency of cancer vaccines is greatly enhanced by the use of adjuvants. The selection of an adjuvant for use with a particular vaccine can have a
20 beneficial effect on the clinical outcome of vaccination. Some vaccines are ineffective in the absence of an adjuvant. Effectiveness of a vaccine may be particularly troublesome when the vaccine is produced from self antigens such as those required for cancer vaccines or
25 other non-infectious disease vaccines. In view of the beneficial effects of adjuvants in vaccine formulations, it is surprising that only one type of adjuvant, aluminum-salt based adjuvants, are currently in wide use in United States-licensed vaccines.

Thus, there exists a need for more and improved immunological adjuvants. The present invention satisfies this need and provides related advantages as well.

SUMMARY OF THE INVENTION

5 The invention provides an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence GAVQNRFNSAIT, or a modification thereof, and having toll-like receptor 5 (TLR5) binding. Methods of inducing an immune response
10 are also provided.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows NF- κ B activation and TNF α production in cells expressing CD4-TLR4 or CD4-TLR5.

15 Figure 2 shows selective induction of TLR5-stimulated activation of NF- κ B by *P. aeruginosa* and *L. monocytogenes* cultures compared to LPS and lipopeptide.

20 Figure 3 shows the purification of a TLR5-stimulating activity from *L. monocytogenes* culture supernatant.

Figure 4 shows the identification by mass spectrometry of flagellin as a TLR5-stimulating activity.

Figure 5 shows that flagellin expression in bacteria reconstitutes TLR5-stimulating activity.

Figure 6 shows systemic induction of IL-6 in wild type mice treated with purified flagellin.

Figure 7 shows a comparison of flagellin amino acid sequences from 22 species of bacteria and a
5 consensus sequence of amino acid residues conserved across species.

DETAILED DESCRIPTION OF THE INVENTION

The invention is directed to flagellin derived peptides that exhibit immunomodulatory activity and to
10 methods of inducing an immune response through activation of toll-like receptor 5 (TLR5). The identification of active flagellin peptides and their corresponding receptor, TLR5, expands the available treatment methods for inducing an immune response. Moreover, the
15 identification of active flagellin peptides and their cognate receptor allows the identification of immunomodulatory compounds.

In one embodiment, the invention is directed to immunomodulatory flagellin peptides that bind to TLR5 and
20 induce a TLR5-mediated activity. The peptides can be used, for example, to effectively stimulate an immune response or ameliorate a pathological condition by administration of immunomodulatory flagellin peptides and combinations of such peptides with antigens and other
25 immunomodulatory molecules. Full length flagellin polypeptides are also used in the methods of the invention to stimulate an immune response. An advantage of the immunomodulatory flagellin peptides of the invention is that they provide the specificity of

flagellin together with the availability of rapid and efficient methods for recombinant and chemical synthesis of peptides. The immunomodulatory flagellin peptides of the invention can therefore be combined with numerous
5 well known modes of administration for the treatment of a wide variety of pathological conditions.

In another embodiment, the invention provides a method of inducing an immune response in an individual by administering a vaccine containing an immunomodulatory
10 flagellin peptide of the invention and an antigen. An immunomodulatory flagellin peptide of the invention functions to stimulate an innate immune response. The innate immune response involves the production of immunomodulatory molecules that beneficially promote the
15 adaptive immune response. The adaptive immune response includes both humoral and cell-mediated immune responses to antigen. Thus, a flagellin peptide functions to boost either or both humoral and cell-mediated immune responses against the antigen. A boost in an immune response
20 causes a general increase in immune system activity that can result in the destruction of foreign or pathologically aberrant cells that otherwise could have escaped the immune response.

As used herein, the term "flagellin" is
25 intended to mean a flagellin polypeptide contained in a variety of Gram-positive or Gram-negative bacterial species. The nucleotide and amino acid sequences of flagellin from 22 bacterial species are depicted in Figure 7. The nucleotide sequences encoding the listed
30 flagellin polypeptides are publically available in the NCBI Genbank database. The flagellin sequences from

these and other species are intended to be encompassed by the term flagellin as used herein. Therefore, the sequence differences between species is included within the meaning of the term.

5 As used herein, the term "peptide" is intended to mean two or more amino acids covalently bonded together. The term "flagellin peptide" is intended to mean a peptide or fragment encoded by a portion of the nucleotide sequence or having a portion of the amino acid
10 sequence which exhibits substantially the same sequence identity to the flagellin sequences as described above and identified in Figure 7 and binds to toll-like receptor 5 (TLR5). For example, a flagellin peptide amino acid sequence is about 65% or greater in sequence
15 identity to a portion of the *S. Typhimurium*1 sequence, GAVQNRFNSAIT, identified as SEQ ID NO:2, encoded by the nucleic acid sequence identified as SEQ ID NO:1. Therefore, flagellin peptides having amino acid substitutions that do not substantially alter TLR5
20 binding are included within the definition of a flagellin peptide. For example, flagellin peptides which contain one or more alanine substitutions and have substantially the same TLR5 binding activity as the flagellin peptide identified as SEQ ID NO:2 are included within the
25 definition of a flagellin peptide. Exemplary flagellin peptides containing alanine substitutions and having substantially the same TLR5 binding activity as the flagellin peptide identified as SEQ ID NO:2 include, for example, GAVANRFNSAIT and GAVQNAFNSAIT. Flagellin
30 peptides consisting of greater than twelve amino acids and having TLR5 binding activity can similarly contain amino acid substitutions, so long as such substituted

peptides retain substantially the same TLR5 binding activity. Examples of such flagellin peptides containing substitutions of various amino acid residues with alanine include ADTRDLGAVQNRFN~~SAIT~~, VDARDLGAVQNRFN~~SAIT~~ and
5 VDTADLGAVQNRFN~~SAIT~~. A flagellin peptide of the invention does not include a full length flagellin polypeptide. A flagellin peptide is intended to include molecules which contain, in whole or in part, non-amide linkages between amino acids, amino acid analogs and mimetics. Similarly,
10 a flagellin peptide also includes cyclic peptides and other conformationally constrained structures. A flagellin peptide of the invention includes polypeptides having several hundred or more amino acid residues and can contain a heterologous amino acid sequence.

15 The term flagellin peptide specifically excludes fragments of flagellin described in Newton et al. Science, 244:70-72 (1989); Kuwajima, G., J. Bacteriol. 170:3305-3309 (1988); McSorley et al., J. Immunol. 164:986-993 (2000); and Samatey et al. J. Struct. Biol. 132:106-111 (2000).
20

As used herein, term "immunomodulatory flagellin peptide," is intended to mean a peptide or fragment having a portion of the amino acid sequence which exhibits substantially the same sequence identity
25 to the flagellin sequences as described above and shown in Figure 7 and binds to toll-like receptor 5 (TLR5). For example, an immunomodulatory flagellin peptide amino acid sequence is about 65% or greater in sequence identity to a portion of the *S. Typhimurium*1 sequence,
30 GAVQNRFN~~SAIT~~, identified as SEQ ID NO:2, encoded by the nucleic acid sequence identified as SEQ ID NO:1.

Therefore, immunomodulatory flagellin peptides having amino acid substitutions that do not substantially alter TLR5 binding are included within the definition of an immunomodulatory flagellin peptide. For example,

5 immunomodulatory flagellin peptides which contain one or more alanine substitutions and have substantially the same TLR5 binding activity as the flagellin peptide identified as SEQ ID NO:2 are included within the definition of a flagellin peptide. Exemplary

10 immunomodulatory flagellin peptides containing alanine substitutions and having substantially the same TLR5 binding activity as the flagellin peptide identified as SEQ ID NO:2 include, for example, GAVANRFNSAIT and GAVQNAFNSAIT. Immunomodulatory flagellin peptides

15 consisting of greater than twelve amino acids and having TLR5 binding activity can similarly contain amino acid substitutions, so long as such substituted peptides retain substantially the same TLR5 binding activity. Examples of such immunomodulatory flagellin peptides

20 containing substitutions of various amino acid residues with alanine include ADTRDLGAVQNRFNSAIT, VDARDLGAVQNRFNSAIT and VDTADLGAVQNRFNSAIT. An immunomodulatory flagellin peptide of the invention does not include a full length flagellin polypeptide. An

25 immunomodulatory flagellin peptide is intended to include molecules which contain, in whole or in part, non-amide linkages between amino acids, amino acid analogs and mimetics. Similarly, an immunomodulatory flagellin peptide also includes cyclic peptides and other

30 conformationally constrained structures. An immunomodulatory flagellin peptide of the invention includes polypeptides having several hundred or more

amino acid residues and can contain a heterologous amino acid sequence.

An immunomodulatory flagellin peptide, polypeptide or modification thereof, of the invention
5 binds to toll-like receptor 5 (TLR5) and induces a TLR5-mediated response. It is understood that minor modifications can be made without destroying the TLR5 binding activity, TLR5-mediated response stimulating activity or immune response modulating activity of an
10 flagellin peptide or polypeptide and that only a portion of the primary structure may be required in order to effect activity. Such modifications are included within the meaning of the terms flagellin polypeptide and flagellin peptide so long as TLR5 binding activity, TLR5
15 response stimulating or immune response stimulating activities are retained. Further, various molecules can be attached to flagellin polypeptides and peptides, including for example, other polypeptides, carbohydrates, nucleic acids or lipids. Such modifications are included
20 within the definition of the term.

Minor modifications of flagellin polypeptide and peptides having at least about the same TLR5 binding activity, TLR5 response stimulating or immune response stimulating activity as the referenced polypeptide or
25 peptide include, for example, conservative substitutions of naturally occurring amino acids and as well as structural alterations which incorporate non-naturally occurring amino acids, amino acid analogs and functional mimetics. For example, a Lysine (Lys) is considered to
30 be a conservative substitution for the amino acid Arg. Similarly, a flagellin peptide containing mimetic

structures having similar charge and spacial arrangements as reference amino acid residues would be considered a modification of the reference polypeptide or peptide so long as the peptide mimetic exhibits at least about the same activity as the reference peptide.

As used herein, the term "amino acid" is intended to mean both naturally occurring and non-naturally occurring amino acids as well as amino acid analogs and mimetics. Naturally occurring amino acids include the 20 (L)-amino acids utilized during protein biosynthesis as well as others such as 4-hydroxyproline, hydroxylysine, desmosine, isodesmosine, homocysteine, citrulline and ornithine, for example. Non-naturally occurring amino acids include, for example, (D)-amino acids, norleucine, norvaline, p-fluorophenylalanine, ethionine and the like. Amino acid analogs include modified forms of naturally and non-naturally occurring amino acids. Such modifications can include, for example, substitution or replacement of chemical groups and moieties on the amino acid or by derivitization of the amino acid. Amino acid mimetics include, for example, organic structures which exhibit functionally similar properties such as charge and charge spacing characteristic of the reference amino acid. For example, an organic structure which mimics Arginine (Arg or R) would have a positive charge moiety located in similar molecular space and having the same degree of mobility as the ϵ -amino group of the side chain of the naturally occurring Arg amino acid. Mimetics also include constrained structures so as to maintain optimal spacing and charge interactions of the amino acid or of the amino acid functional groups. Those skilled in the art know or

can determine what structures constitute functionally equivalent amino acid analogs and amino acid mimetics.

Specific examples of amino acid analogs and mimetics can be found described in, for example, Roberts and Vellaccio, The Peptides: Analysis, Synthesis, Biology, Eds. Gross and Meinhofer, Vol. 5, p. 341, Academic Press, Inc., New York, New York (1983), the entire volume of which is incorporated herein by reference. Other examples include peralkylated amino acids, particularly permethylated amino acids. See, for example, Combinatorial Chemistry, Eds. Wilson and Czarnik, Ch. 11, p. 235, John Wiley & Sons Inc., New York, New York (1997), the entire book of which is incorporated herein by reference. Yet other examples include amino acids whose amide portion (and, therefore, the amide backbone of the resulting peptide) has been replaced, for example, by a sugar ring, steroid, benzodiazepine or carbo cycle. See, for instance, Burger's Medicinal Chemistry and Drug Discovery, Ed. Manfred E. Wolff, Ch. 15, pp. 619-620, John Wiley & Sons Inc., New York, New York (1995), the entire book of which is incorporated herein by reference. Methods for synthesizing peptides, polypeptides, peptidomimetics and proteins are well known in the art (see, for example, U.S. Patent No. 5,420,109; M. Bodanzsky, Principles of Peptide Synthesis (1st ed. & 2d rev. ed.), Springer-Verlag, New York, New York (1984 & 1993), see Chapter 7; Stewart and Young, Solid Phase Peptide Synthesis, (2d ed.), Pierce Chemical Co., Rockford, Illinois (1984), each of which is incorporated herein by reference).

As used herein, the term "immune response" is intended to mean to a measurable or observable reaction to an antigen or immunomodulatory molecule mediated by one or more cells of the immune system. An immune
5 response begins with an antigen or immunomodulatory molecule binding to an immune system cell and terminates with destruction of antigen and cells containing antigen or alteration in immune cell function. A reaction to an antigen or immunomodulatory molecule is mediated by many
10 cell types, including a cell that initially binds to an antigen or immunomodulatory molecule and cells that participate in mediating an innate, humoral, cell-mediated immune response. An innate immune response involves binding of pathogen-associated molecular
15 patterns (PAMPs) to cell surface receptors, such as toll-like receptors. Activation of toll-like receptors in response to PAMPs leads to the production of immunomodulatory molecules, such as cytokines and co-stimulatory molecules, that induce an immune response. A
20 humoral response involves interaction of B cells with antigen and B cell differentiation into antibody-secreting cells. A cell-mediated response involves various subpopulations of T cells that recognize antigen presented on self-cells, including helper T cells
25 that respond to antigen by producing cytokines and cytotoxic T cells that respond to antigen by developing into cytotoxic T lymphocytes, which mediate killing of altered self-cells. The term immune response includes measurable or observable reactions produced by any cell
30 type that participates in the processes through which immune system cells are activated and antigen containing cells are destroyed. Such measurable reactions include,

for example, production of immunomodulatory molecules, migration and proliferation.

An "immunomodulatory molecule" is a molecule that alters an immune response. An immunomodulatory molecule can be, for example, a compound, such as an organic chemical; a polypeptide, such as an antibody or cytokine; a nucleic acid, such as a DNA or RNA molecule; or any other type of molecule that alters an immune response. An immunomodulatory molecule can alter an immune response by directly or indirectly altering an activity of a cell that mediates an immune response. An immunomodulatory molecule can act directly on an immune system cell, for example, by binding to a cell surface receptor and stimulating or inhibiting proliferation, differentiation, or expression, secretion or receptor binding of immune system regulatory molecules such as co-stimulatory receptors and ligands, cytokines, and chemokines. Examples of naturally occurring molecules that act directly on immune system cells to alter an immune response include PAMPs, cytokines, chemokines and growth factors. Other examples of molecules that act directly on immune system cells to alter an immune response include molecules that alter receptor functions, such as antibodies to receptors, soluble cytokine receptors, receptor agonists and antagonists, molecules that alter the production of immunomodulatory molecules, such as inhibitors of converting enzymes and molecules involved in the intracellular transport and secretion of immunomodulatory molecules.

An immunomodulatory molecule can indirectly alter the activity of a particular immune system cell by

altering the amount or activity of a molecule that regulates a cellular activity of the cell. For example, a cytokine, chemokine, or growth factor produced by an immune system cell, such as a macrophage, can stimulate or inhibit various cellular activities of B and T lymphocytes. Immune cell functions that can be stimulated or inhibited by an immunomodulatory molecule include, for example, immune cell activation, co-activation, proliferation, production of cytokines, cellular interactions and migration. An immunomodulatory molecule can therefore act on a variety of immune cell types and can alter a variety of cellular functions. An immunomodulatory flagellin peptide, polypeptide or modifications thereof used in the methods of the invention are examples of immunomodulatory molecules useful for inducing an immune response, for example, by binding to TLR5 and inducing a TLR5-mediated increase in macrophage production of $\text{TNF}\alpha$, IL-1 and IL-6. The flagellin polypeptides, peptides and modifications thereof, are also useful for indirectly inducing an immune response because immunomodulatory molecules produced by a TLR5-expressing cell in response to flagellin will alter the activities of immune system cells that respond to the particular immunomodulatory molecules produced.

An immunomodulatory molecule can mediate an immune response that is specific for a target antigen or nonspecific. A specific immunomodulatory molecule alters an immune response to a particular target antigen. Examples of specific immunomodulatory molecules include monoclonal antibodies, including naked monoclonal antibodies, drug-, toxin- or radioactive

compound-conjugated monoclonal antibodies, and ADCC targeting molecules. Such immunomodulatory molecules stimulate an immune response by binding to antigens and targeting cells for destruction. An immunomodulatory molecule can be used to suppress an immune response to an antigen. For example, a tolerogenizing molecule can be used to suppress an immune response to a self-antigen.

Nonspecific immunomodulatory molecules stimulate or inhibit the immune system in a general manner through various mechanisms that can include, for example, stimulating or suppressing cellular activities of immune system cells. Nonspecific immunomodulatory molecules useful for stimulating an immune responses include, for example, agents that stimulate immune cell proliferation, immune cell activation and production of cytokines and co-stimulatory molecules. Well known immunomodulatory molecules that stimulate an immune response are, for example, interleukins, interferons, levamisole and keyhole limpet hemocyanin. Nonspecific immunomodulatory molecules useful for suppressing immune responses include, for example, agents that inhibit cytokines synthesis or processing, specific cytokine receptor blocking reagents such as soluble receptors and receptor antagonists, and cytokines that down-regulate or inhibit the production of other immunomodulatory molecules. Well known immunomodulatory molecules for suppressing an immune response include, for example, cyclosporin, rapamycin, tacrolimus, azathioprine, cyclophosphamide and methotrexate.

Immunomodulatory molecules can be contained in a mixture of molecules, including a natural or man-made

composition of molecules. Exemplary natural compositions of immunomodulatory compounds include, for example, those contained in an organism such as Bacille Calmette-Guerin (BCM) or *Corynebacterium parvum*. Exemplary man-made
5 compositions of immunomodulatory molecules include, for example, QS-21, DETOX and incomplete Freund's adjuvant.

As used herein, the term "adjuvant" when used in reference to a vaccine, is intended to mean a substance that acts generally to accelerate, prolong, or
10 enhance the quality of specific immune responses to a vaccine antigen. An adjuvant can advantageously reduce the number of immunizations or the amount of antigen required for protective immunization.

15 As used herein, the term "antigen-specific immune response" is intended to mean a reaction of one or more cells of the immune system to a particular antigen that is not substantially cross-reactive with other antigens.

20 As used herein, the term "antigen" is intended to mean a molecule which induces an immune response. An antigen can be a crude mixture of molecules, such as a cell, or one or more isolated molecules. Examples of crude antigens include attenuated organisms, inactivated
25 organisms, viral particles and tumor cells. Examples of isolated antigens include a polypeptide, lipoprotein, glycoprotein, lipid, anti-idiotypic antibody, toxoid, polysaccharide, capsular polysaccharide and nucleic acid. Such isolated antigens can be naturally occurring,
30 recombinantly produced, or synthesized. Exemplary naturally occurring antigens include purified microbial

macromolecules. Exemplary recombinantly produced antigens include cloned microbial and tumor cell antigens. Exemplary synthesized antigens include synthetic peptides and nucleic acids.

5 As used herein, the term "vaccine" is intended to mean a compound or formulation which, when administered to an individual, stimulates an immune response against an antigen. A vaccine is useful for preventing or ameliorating a pathological condition that
10 will respond favorably to immune response modulation. A vaccine can contain isolated or crude antigen, and can contain one or more antigens. A vaccine can contain one or more adjuvants.

 As used herein, the term "immunogenic amount"
15 is intended to mean an amount of an immunomodulatory flagellin polypeptide, peptide or modifications thereof, or combinations thereof with one or more molecules, such as an antigen or other immunomodulatory molecule, required to effect an immune response. The dosage of an
20 immunomodulatory flagellin polypeptide, peptide, or modifications thereof, independently or in combination with one or more molecules, will depend, for example, on the pathological condition to be treated, the weight and condition of the individual and previous or concurrent
25 therapies. The appropriate amount considered to be an immunogenic dose for a particular application of the method can be determined by those skilled in the art, using the guidance provided herein. For example, the amount can be extrapolated from *in vitro* or *in vivo*
30 assays as described below. Those skilled in the art will understand that the condition of the patient needs to be

monitored through the course of therapy and that the amount of the composition that is administered can be adjusted according to patient response to therapy.

The term "pathologically aberrant cell" is intended to mean a cell that is altered from a normal physiological or cellular state. Such alteration can be due to changes in physiology or phenotype associated with a disease or abnormal condition of a mammalian cell or tissue. Pathologically aberrant cells include cells lacking normal control of cellular functions, such as growth, differentiation, and apoptosis, resulting in altered gene and protein expression. Cells that lack normal growth control proliferate in the absence of appropriate growth signals, resulting in damage in structure or function of surrounding tissues. Cells that lack normal differentiation undergo inappropriate phenotypic or physiological changes that do not normally characterize the cell type, resulting in damage in structure and function of surrounding tissues. Cells that lack normal apoptosis fail to undergo, or inappropriately undergo the process of cell death, resulting in damage in structure or function of surrounding tissues. Altered protein expression is an example of a phenotype change that renders such cells distinguishable from normal. For example, increased or decreased expression of a polypeptide normally expressed on a cell, expression of a mutated polypeptide and expression of a polypeptide not normally expressed on a cell are phenotypic changes that can alter a cell from normal. Examples of pathologically aberrant cells include tumor cells and degenerating cells.

As used herein, the term "pathological condition" is intended to mean a disease, abnormal condition or injury of a mammalian cell or tissue. Such pathological conditions include, for example,

5 hyperproliferative and unregulated neoplastic cell growth, degenerative conditions, inflammatory diseases, autoimmune diseases and infectious diseases. Pathological conditions characterized by excessive or unregulated cell growth include, for example,

10 hyperplasia, cancer, autoimmune disease and infectious disease. Hyperplastic and cancer cells proliferate in an unregulated manner, causing destruction of tissues and organs. Specific examples of hyperplasias include benign prostatic hyperplasia and endometrial hyperplasia.

15 Specific examples of cancer include prostate, breast, ovary, lung, uterus, brain and skin cancers. Abnormal cellular growth can also result from infectious diseases in which foreign organisms cause excessive growth. For example, human papilloma viruses can cause abnormal

20 growth of skin cells. The growth of cells infected by a pathogen is abnormal due to the alteration of the normal condition of a cell resulting from the presence of a foreign organism. Specific examples of infectious diseases include DNA and RNA viral diseases, bacterial

25 diseases, parasitic diseases. Similarly, the growth of cells mediating autoimmune and inflammatory diseases are aberrantly regulated which results in, for example, the continued proliferation and activation of immune mechanisms with the destruction of tissues and organs.

30 Specific examples of autoimmune diseases include, for example, rheumatoid arthritis and systemic lupus erythematosus. Specific examples of degenerative disease include osteoarthritis and Alzheimer's disease.

By specific mention of the above categories of pathological conditions, those skilled in the art will understand that such terms include all classes and types of these pathological conditions. For example, the term
5 cancer is intended to include all known cancers, whether characterized as malignant, benign, soft tissue or solid tumor. Similarly, the terms infectious diseases, degenerative diseases, autoimmune diseases and
10 inflammatory diseases are intended to include all classes and types of these pathological conditions. Those skilled in the art will know the various classes and types of proliferative, infectious, autoimmune and inflammatory diseases.

As used herein the term "toll-like receptor 5"
15 or "TLR5" is intended to mean a toll-like receptor 5 of any species, such as the murine and human polypeptides containing the amino acid sequences set forth as SEQ ID NOS:4 and 6, respectively, encoded by the nucleic acid sequence identified as SEQ ID NOS:3 and 5, respectively.
20 A TLR5 is activated upon binding to flagellin, an immunomodulatory flagellin peptide, or modifications thereof, and other TLR5 agonists. Upon activation, a TLR5 induces a cellular response by transducing an intracellular signal that is propagated through a series
25 of signaling molecules from the cell surface to the nucleus. For example, the intracellular domain of TLR5 recruits an adaptor protein, MyD88, which recruits the serine kinase IRAK. IRAK forms a complex with TRAF6, which then interacts with various molecules that
30 participate in transducing the TLR signal. These molecules and other TLR5 signal transduction pathway components stimulate the activity of transcription

factors, such as fos, jun and NF- κ B, and the corresponding induction of gene products of fos-, jun- and NF- κ B-regulated genes, such as, for example, TNF α , IL-1 and IL-6. The activities of signaling molecules that mediate the TLR5 signal, as well as molecules produced as a result of TLR5 activation are TLR5 activities that can be observed or measured. Therefore, a TLR5 activity includes binding to a flagellin polypeptide, immunomodulatory flagellin peptide, or a modification thereof, recruitment of intracellular signaling molecules, as well as downstream events resulting from TLR5 activation, such as transcription factor activation and production of immunomodulatory molecules. A TLR5 cellular response mediates an innate immune system response in an animal because cytokines released by TLR5-expressing cells regulate other immune system cells to promote an immune response in an animal. Therefore, as used herein the term "TLR5-mediated response" is intended to mean the ability of a flagellin polypeptide, immunomodulatory peptide or modification thereof to induce a TLR5-mediated cellular response. Exemplary TLR5-mediated cellular responses include activation of transcription factors such as fos, jun and NF- κ B, production of cytokines such as IL-1, IL-6 and TNF α , and the stimulation of an immune response in an animal.

A TLR5 also encompasses polypeptides containing minor modifications of a native TLR5, and fragments of a full-length native TLR5, so long as the modified polypeptide or fragment retains one or more biological activities of a native TLR5, such as the abilities to

stimulate NF- κ B activity, stimulate the production of cytokines such as TNF α , IL-1, and IL-6 and stimulate an immune response in response to TLR5 binding to flagellin polypeptide, immunomodulatory peptide or modifications thereof. A modification of a TLR5 can include additions, deletions, or substitutions of amino acids, so long as a biological activity of a native TLR5 is retained. For example, a modification can serve to alter the stability or activity the polypeptide, or to facilitate its purification. Modifications of polypeptides as described above in reference to flagellin polypeptides and peptides are applicable to TLR5 polypeptides of the invention. A "fragment" of a TLR5 is intended to mean a portion of a TLR5 that retains at least about the same activity as a native TLR5.

As used herein, the term "TLR5 agonist" refers to a compound that selectively activates or increases normal signal transduction through TLR5. As used herein, the term "TLR5 antagonist" refers to a compound that selectively inhibits or decreases normal signal transduction through TLR5. A TLR5 agonist or antagonist can alter normal signal transduction through TLR5 indirectly, for example, by modifying or altering the native conformation of TLR5 or a TLR5 ligand. For therapeutic applications, a TLR5 agonist or antagonist has an EC₅₀ of less than about 10^{-7} M, such as less than 10^{-8} M and less than 10^{-9} M, although a TLR5 agonist with a higher EC₅₀ can be therapeutically useful. As used herein, the term "TLR5 ligand" refers to a compound that binds a TLR5 polypeptide with high affinity. A TLR5 ligand can further be an agonist or antagonist of TLR5,

as described above, or can be a compound having little or no effect on TLR5 signaling.

As used herein, the term "detectably labeled" refers to derivitization with, or conjugation to, a moiety that is detectable by an analytical or qualitative method. A detectable moiety can be, for example, a radioisotope, such as ^{14}C , ^{131}I , ^{32}P or ^3H , fluorochrome, ferromagnetic substance, or luminescent substance.

As used herein the term "ADCC targeting molecule" is intended to mean an antigen binding protein containing a Fc receptor binding domain capable of inducing antibody-dependent cell cytotoxicity (ADCC). An ADCC targeting molecule can also contain other domains that augment induction of ADCC. The flagellin polypeptides and peptides, immunomodulatory peptides, and modifications described herein, can be domains of an ADCC targeting molecule that augment induction of ADCC. The ADCC targeting molecule can include multiple valencies for either or both the antigen binding domain or the Fc receptor binding domain. Additionally, an ADCC targeting molecule also can have multiple different antigen binding domains combined with a single or multiple copies of an Fc receptor binding domain or combined with different Fc receptor binding domains. The antigen binding domain or domains can be derived from essentially any molecule that has selective or specific binding activity to a target antigen so long as it can be fused or attached to one or more Fc receptor binding domains while still maintaining antigen binding activity. The Fc receptor binding domain can be derived from an antibody constant region of, for example, the IgG class, including subclasses IgG1, IgG3

and IgG4. Such Fc receptor binding domains can be used in their native form or the amino acid sequence can be modified so as to enhance or optimize the Fc receptor binding or ADCC activity. Moreover, the Fc receptor binding domains can be derived from constant regions which recognize either stimulatory or inhibitory Fc receptors. The Fc receptor binding domain is located within the hinge region of an antibody constant region where the cognate receptors bound by this domain include, for example, the Fc RI, Fc RIIA and Fc RIII. Therefore, ADCC targeting molecules include, for example, antibodies selective for a target antigen and functional variants thereof as well as fusion proteins and chemical conjugates containing both an antigen binding domain and a Fc receptor binding domain in functionally active forms. ADCC targeting molecules and the use of ADCC targeting molecules in the treatment of disease are described in detail in U.S. Patent Application 09/618,176, which is incorporated herein by reference.

The term "about" when used in reference to a particular activity or measurement is intended to refer to the referenced activity or measurement as being within a range values encompassing the referenced value and within accepted standards of a credible assay within the art, or within accepted statistical variance of a credible assay within the art.

As used herein, the term "substantially" or "substantially the same" when used in reference to an amino acid sequence is intended to mean that the amino acid sequence shows a considerable degree, amount or extent of sequence identity when compared to the

reference sequence. Such considerable degree, amount or extent of identity is further considered to be significant and meaningful and therefore exhibit characteristics which are definitively recognizable or known as being derived from or related to flagellin. For example, an amino acid sequence which is substantially the same amino acid sequence as an flagellin peptide, including fragments thereof, refers to a sequence which exhibits characteristics that are definitively known or recognizable as being sufficiently related to flagellin so as to fall within the classification of flagellin sequences as defined above. Minor modifications thereof are included so long as they are recognizable as an flagellin sequence as defined above.

As used herein, the term "individual" is intended to mean any animal in which an immune response can be induced by a flagellin polypeptide, peptide or modifications thereof including a human, non-human primate, cow, pig, chicken, rabbit, ferret, rat or mouse.

An immunomodulatory flagellin polypeptide, peptide or modifications thereof can be used to induce an immune response in an individual having a pathological condition, promoting the individual's own immune system to function more effectively and thereby ameliorate the pathological condition. An individual's immune system may not recognize cancer cells and other types of pathologically aberrant cells as foreign because the particular antigens are not different enough from those of normal cells to cause an immune reaction. In addition, the immune system may recognize cancer cells, but induce a response insufficient to destroy the cancer.

By stimulating an innate immune response, immunomodulatory flagellin peptide, polypeptide or modification thereof, promote humoral and cell-mediated responses to antigens on foreign cells or pathologically aberrant cells, such as cancer cells. Administered independently or in combination with an antigen, such as a tumor antigen, a flagellin polypeptide, peptide or modification thereof, can be used to boost the immune system's recognition of cancer cells and other pathologically aberrant cells, and target such cells for destruction.

Flagellin is a pathogen-associated molecular pattern (PAMP) recognized by toll-like receptor 5 (TRL5). Toll-like receptor 5 is a member of a family of at least 10 receptors involved in mediated the innate immune response. Toll-like receptors recognize PAMPs that distinguish infectious agents from self and mediating the production of immunomodulatory molecules, such as cytokines, necessary for the development of effective adaptive immunity (Aderem, A and Ulevitch, R.J. Nature 406:782-787 (2000) and Brightbill, H.D., Immunology 101: 1-10 (2000)). Members of the toll-like receptor family recognize a variety of antigen types and can discriminate between pathogens. For example, TLR2 recognizes various fungal, Gram-positive, and mycobacterial components, TLR4 recognizes the Gram-negative product lipopolysaccharide (LPS), and TLR9 recognizes nucleic acids such as CpG repeats in bacterial DNA. TLR5 has now been identified as a receptor for bacterial flagellin.

Flagellin induces an innate immune response by binding to and activating TLR5. Activation of TLR5 by

binding to flagellin induces the production of immunomodulatory molecules, such as cytokines and co-stimulatory molecules, by a TLR5-expressing cell. For example, activation of TLR5 in macrophages results in the
5 expression of the cytokines TNF α , IL-1 and IL-6. These cytokines directly and indirectly alter the activities of immune system cells that participate in both humoral (TH2) and cell-mediated (TH1) adaptive immune responses. In this manner, an immunomodulatory flagellin peptide,
10 polypeptide or modification thereof, acts as an adjuvant to stimulate a general immune response.

Altering the balance of TH1- versus TH2-associated cytokines can be used to favorably alter an
15 immune response to treat certain diseases. For example, in the use of cancer vaccines, it can be favorable to induce both TH1 and TH2 responses (Herlyn and Birebent, Ann. Med., 31:66-78, (1999)). Different sets of cytokines orchestrate TH1 and TH2 immune responses. For
20 example, TH1 immune responses are associated with the cytokines IL-2, IFN- γ , and TNF α while TH2 immune responses are associated with the cytokines IL-4, IL-5, IL-6 and IL-10. TLR5 stimulates the production of cytokines associated with both TH1- and TH2-associated
25 cytokines. For example, TNF α is associated with the stimulation of a TH1 type immune response (Ahlers, JD et al. J. Immunol, 158:3947-58 (1997)), and IL-6 is associated with the stimulation of a TH2 type response (Steidler, L. et al. Infect. Immun., 66:3183-9, (1998)).
30 Therefore, an immunomodulatory flagellin peptide, polypeptide or modification thereof, can be used to advantageously elicit TH1 and TH2 type immune responses.

An immunomodulatory flagellin peptide, polypeptide or modification thereof can also be used to generally alter the particular cytokines involved in an immune response in an individual. Alterations from
5 normal levels of cytokines are observed in many disease states. For this reason, it can be desirable to increase or decrease the amounts or activities of specific cytokines involved in particular pathological conditions. The cytokines produced in response to TLR5 activation can
10 both stimulate and down-regulate the production of other cytokines. Therefore, an immunomodulatory flagellin peptide, polypeptide or modification thereof, or combination of a flagellin molecule with an immunomodulatory molecule or antigen can be used to alter
15 levels of cytokines associated with a pathological condition. For example, an immunomodulatory flagellin peptide can increase TLR5-expressing macrophage production of TNF α , IL-1 and IL-6. TNF α and IL-1 generally function as pro-inflammatory cytokines. IL-6
20 generally functions as an anti-inflammatory cytokine and induces a variety of anti-inflammatory activities in immune system cells. For example, IL-6 stimulates the production of many anti-inflammatory anti-proteases. Those skilled in the art will be able to determine if a
25 pathological condition in an individual could be ameliorated by inducing TLR5-stimulated cytokine production and will be able to determine appropriate combinations of flagellin and immunomodulatory molecules suitable for inducing a beneficial immune response.

30 The invention provides an immunomodulatory flagellin peptide comprising at least about 10 amino acids of substantially the amino acid sequence

GAVQNRFNSAIT (SEQ ID NO:2), or a modification thereof, that binds to toll-like receptor 5 (TLR5).

The flagellin peptide identified by SEQ ID NO:2 is a peptide of *S. Typhimurium*1 flagellin which is encoded by the nucleic acid sequence identified by SEQ ID NO:1. A flagellin peptide of the invention also includes peptides from other bacterial species, such as *H. Pylori*, *V. Cholera*, *S. marcesens*, *S. flexneri*, *T. Pallidum*, *L. pneumophila*, *B burgdorferi*, *C. difficile*, *R. meliloti*,
10 *A. tumefaciens*, *R. lupini*, *B. clarridgeiae*, *P. Mirabilis*, *B. subtilus*, *L. monocytogenes*, *P. aeruginosa* and *E. coli*, which contain amino acid sequences having 21-71% identity over the 12 amino acid sequence of SEQ ID NO:2. Thus, a flagellin peptide of the invention can have greater than
15 about 65% identity, such as greater than about 75%, greater than about 85%, greater than about 95%, greater than about 98% identity with the peptide identified by SEQ ID NO:2.

20 A flagellin peptide of the invention is derived from a conserved region of a flagellin polypeptide. Conserved regions of flagellin are well known in the art and have been described, for example, in Mimori-Kiyosue, et al., J. Mol. Biol. 270:222-237, (1997). Whereas
25 T cell receptors which mediate the adaptive immune response recognize random portions of antigen amino acid sequences, toll-like receptors recognize conserved portions of antigen amino acid sequences. Therefore, the flagellin peptides of the invention and immunomodulatory
30 flagellin peptides used in the methods of the invention contain amino acid sequences derived from conserved regions of flagellin.

A flagellin peptide of the invention excludes a portion of flagellin described in Newton et al. (supra, 1989), which consists of an *S. meunchen* flagellin fragment containing a deletion of amino acids 207-223, portions of *E. coli* (strain K12) flagellin described in Kuwaijima et al. (supra, 1998), which consist of *E. coli* flagellin fragments containing deletions of amino acids 239-254, 259-278, 237-262, 194-379, 201-318, 218-326, 211-347, 210-299, 245-301, and 220-299, a portion of flagellin described in Samatey et al. (supra, 2000), which consists of an *S. typhimurium* flagellin fragment lacking 52 N-terminal amino acid residues and lacking 44 C-terminal amino acid residues, and portions of flagellin described in McSorley et al. (supra, 2000) which consist of *S. typhimurium* flagellin fragments having the following amino acid sequences: RSDLGAVQNRFNSAI, DLGAVQNRFNSAITN, GAVQNRFNSAITNLG AND VQNRFNSAITNLGNT.

An immunomodulatory flagellin peptide of the invention can contain a heterologous amino acid sequence that imparts structural or functional characteristics onto the flagellin peptide. For example, chimeric flagellin peptides or modifications can be used to impart a targeting function. Targeting of a flagellin peptide or modification to a particular site, such as a mucosal surface for example, confers additional therapeutic advantage of inducing an immune response at a site of pathological condition or a site favored for inducing an antigen-specific immune response, for example by a vaccine. Further, chimeric flagellin peptides can include a sequence that facilitates detection, purification or enhances immunomodulatory activity of the flagellin peptide. A flagellin peptide can be contained,

for example, in an ADCC targeting molecule used to treat a pathological condition. A flagellin peptide can augment the effectiveness of an ADCC targeting molecule by, for example, stimulating an innate immune response through TLR5, such as the induction of cytokines such as TNF α , IL-1 and IL-6. Similarly, a flagellin peptide can contain amino acid sequences of a variety of antigen polypeptides, such as those described above in reference to antigens contained in vaccines used in the methods of the invention. A chimeric flagellin peptide containing amino acid sequences of an antigen or containing an antigenic molecule such as a carbohydrate, nucleic acid, or lipid, can be used analogously to a vaccine, as described above, as well as in a vaccine formulation, to induce an immune response in an individual. As such, a chimeric flagellin peptide can be a vaccine that induces both innate and adaptive immune system responses.

An immunomodulatory flagellin peptide of the invention can be prepared by a variety of methods well-known in the art, for example, by recombinant expression systems described below, and biochemical purification methods described below, as well as by synthetic methods well known in the art. Methods for recombinant expression and purification of polypeptides in various host organisms are described, for example, in Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, New York (1992) and in Ansubel et al., Current Protocols in Molecular Biology, John Wiley and Sons, Baltimore, MD (1998), both of which are incorporated herein by reference. Similarly, flagellin peptide modifications can be generated using recombinant mutagenesis, such as site directed

mutagenesis and PCR mutagenesis, and expression of the flagellin peptide modification. Numerous methods of constructing, modifying, expressing and purifying peptides are known to those skilled in the art. A
5 specific example of a method for purifying flagellin is described below in Example III. The choice of recombinant methods, expression and purification systems will be known by those skilled in the art and will depend on the user and the particular application for the
10 immunomodulatory flagellin peptide or modification thereof.

A flagellin peptide of the invention induces an innate immune response in an individual by binding to an stimulating TLR5. Therefore, the invention provides
15 methods for inducing an immune response in an individual having a pathological condition that can be ameliorated by immune system activity. The methods involve administering an immunomodulatory flagellin peptide or modification thereof to induce an immune response,
20 administering a combination of an immunomodulatory flagellin peptide and an antigen to induce an antigen-specific immune response, and administering a combination of an immunomodulatory flagellin peptide and an immunomodulatory molecule to modulate an immune response.
25 The selection of a particular method for inducing an immune response will depend on the particular pathological condition to be ameliorated or prevented in an individual. As described herein, the methods are applicable to a wide variety of pathological conditions.
30 Those skilled in the art will be able to determine if an immune response can be beneficially modulated by administering an immunomodulatory flagellin peptide or

combination thereof with an antigen or immunomodulatory molecule.

The invention provides method of inducing an antigen-specific immune response in an individual. The method involves administering to an individual an immunogenic amount of a vaccine, comprising an antigen and an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof.

As an adjuvant in a vaccine formulation, the immunomodulatory flagellin peptides of the invention can contribute to the effectiveness of the vaccine by, for example, enhancing the immunogenicity of weaker antigens such as highly purified or recombinant antigens, reducing the amount of antigen required for an immune response, reducing the frequency of immunization required to provide protective immunity, improve the efficacy of vaccines in individuals with reduced or weakened immune responses, such as newborns, the aged, and immunocompromised individuals, and enhance the immunity at a target tissue, such as mucosal immunity, or promote cell-mediated or humoral immunity by eliciting a particular cytokine profile. An immunomodulatory flagellin peptide, polypeptide or modification thereof induces an innate immune response through activation of TLR5. The innate immune response increases the immune response to an antigen by stimulating the adaptive immune response. Therefore, a combination of an immunomodulatory flagellin peptide, polypeptide or modification thereof with one or more antigens provides

an effective vaccine for inducing an immune response in an individual.

The methods of the invention for inducing an antigen-specific immune response can be used to treat
5 individuals having a variety of pathological conditions. For example, cancer vaccines have been used effectively for treating melanoma and breast cancers. Vaccines have been used for treatment of inflammatory diseases such as asthma (Scanga C.B and Le Gros, G., Drugs 59(6), 1217-
10 1221 (2000)), infectious diseases of pathogenic bacteria such as *H. pylori*, pathogenic viruses such as human papilloma virus and HIV (Sutton P. and Lee, A, Aliment Pharmacol. 14:1107-1118 (2000)), protozoa, autoimmune diseases such as diabetes (von Herrath and Whitton, Ann.
15 Med. 32:285-292 (2000)) and degenerative diseases such as Alzheimer's disease (Youngkin, S.G., Nat. Med., 7(1):18-19 (2001)). Therefore, a vaccine used in the methods of the invention for inducing an antigen-specific immune response can be administered to an individual for
20 treatment of a variety of pathological conditions, including proliferative disease, infectious disease, inflammatory disease and degenerative disease.

A variety of antigens can be used in combination with an immunomodulatory flagellin peptide of
25 the invention for preparing a vaccine. Microorganisms such as viruses, bacteria and parasites contain substances that are not normally present in the body. These substances can be used as antigens to produce an immune response to destroy both the antigen and cells
30 containing the antigen, such as a bacterial cell or cancer cell.

For example, isolated or crude antigens of microbial pathogens can be used in vaccines to treat infectious disease; isolated or crude tumor cell antigens can be used in vaccines to treat cancer; isolated or
5 crude antigens known to be associated with a pathologically aberrant cell can be used to treat a variety of diseases in which it is beneficial to target particular cells for destruction.

A variety of substances can be used as antigens
10 in a vaccine compound or formulation. For example, attenuated and inactivated viral and bacterial pathogens, purified macromolecules, polysaccharides, toxoids, recombinant antigens, organisms containing a foreign gene from a pathogen, synthetic peptides, polynucleic acids,
15 antibodies and tumor cells can be used to prepare a vaccine useful for treating a pathological condition. Therefore, an immunomodulatory flagellin peptide of the invention can be combined with a wide variety of antigens to produce a vaccine useful for inducing an immune
20 response in an individual. Those skilled in the art will be able to select an antigen appropriate for treating a particular pathological condition and will know how to determine whether a crude or isolated antigen is favored in a particular vaccine formulation.

25 An isolated antigen can be prepared using a variety of methods well known in the art. A gene encoding any immunogenic polypeptide can be isolated and cloned, for example, in bacterial, yeast, insect, reptile or mammalian cells using recombinant methods well known
30 in the art and described, for example in Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring

Harbor Laboratory, New York (1992) and in Ansubel et al.,
Current Protocols in Molecular Biology, John Wiley and
Sons, Baltimore, MD (1998). A number of genes encoding
surface antigens from viral, bacterial and protozoan
5 pathogens have been successfully cloned, expressed and
used as antigens for vaccine development. For example,
the major surface antigen of hepatitis B virus, HbsAg,
the β subunit of cholera toxin, the enterotoxin of
E. coli, the circumsporozoite protein of the malaria
10 parasite, and a glycoprotein membrane antigen from
Epstein-Barr virus, as well as tumor cell antigens, have
been expressed in various well known vector/host systems,
purified and used in vaccines. An immunomodulatory
flagellin peptide, polypeptide or modification thereof
15 induces an innate immune response through TLR5 that can
beneficially enhance an immune response to a recombinant
antigen.

A pathologically aberrant cell to be used in a
vaccine can be obtained from any source such as one or
20 more individuals having a pathological condition or
ex vivo or in vitro cultured cells obtained from one or
more such individuals, including a specific individual to
be treated with the resulting vaccine.

Those skilled in the art will be able to
25 determine if a vaccine compound or formulation induces an
innate, humoral, cell-mediated, or any combination of
these types of immune response, as methods for
characterizing these immune responses are well known in
the art. For example, the ability of a vaccine compound
30 or formulation to induce an innate immune response
through TLR5 can be determined using methods described

herein as well as other methods. Such methods for detecting an innate immune response can be generally performed within hours of vaccine administration. The ability of a vaccine compound or formulation to induce a humoral response can be determined by measuring the titer of antigen-specific antibodies in an animal primed with the vaccine and boosted with the antigen, or determining the presence of antibodies cross-reactive with an antigen by ELISA, Western blotting or other well-known methods. Cell-mediated immune responses can be determined, for example, by measuring cytotoxic T cell response to antigen using a variety of methods well known in the art. Methods of detecting humoral and cell-mediated immune responses can be generally performed days or weeks after vaccine administration.

A combination of an antigen or immunomodulatory molecule and an immunomodulatory flagellin peptide, polypeptide or modification thereof, can be tested in a variety of preclinical toxicological and safety studies well known in the art. For example, such a combination can be evaluated in an animal model in which the antigen has been found to be immunogenic and that can be reproducibly immunized by the same route proposed for human clinical testing. A combination of an antigen or immunomodulatory molecule and an immunomodulatory flagellin peptide or modification thereof can be tested, for example, by an approach set forth by the Center for Biologics Evaluation and Research/Food and Drug Administration and National Institute of Allergy and Infectious Diseases (Goldenthal, KL et al. AID Res Hum Retroviruses, 9:S45-9 (1993)).

Those skilled in the art will know how to determine for a particular combination of antigen or immunomodulatory molecule and immunomodulatory flagellin polypeptide modification thereof, the appropriate antigen
5 payload, route of immunization, volume of dose, purity of antigen, and vaccination regimen useful to treat a particular pathological condition in a particular animal species.

10 The invention provides a method of inducing a TLR5-mediated response. The method involves administering to a TLR5-containing cell an effective amount of an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino
15 acid sequence of SEQ ID NO:2, or a modification thereof.

A TLR5-mediated response can be assessed in a cell or animal because TLR5 stimulates cellular activities that stimulate the immune response that occurs
20 in an animal. For example, flagellin binding to TLR5 induces cellular events such as an increase in the amount or activity of cytokines, such as TNF α , IL-1 and IL-6. These cytokines in turn regulate the activities of immune system cells. Therefore a TLR5-mediated response can be
25 determined by examining an immune responses in an animal and by observing particular immune system cell activities. Determination of immune responses in an animal is discussed below. Determination of immune system cell activities can be performed, for example, by
30 observing or measuring the amount of activity of immunomodulatory molecules produced by specific types of immune cells. Cytokine production by macrophages is an exemplary immune cell activity that can be conveniently

measured using methods well known in the art and those described herein. A biological activity of a cytokine can also be assessed using methods well known in the art. TNF α activities include, for example, inducing the

5 production of IL-1 and IL-6, activation of neutrophils and endothelial cells in inflammation, inducing acute phase reactants in liver, inducing fever. IL-1 activities include, for example, activating of endothelial cells in inflammation and coagulation,

10 inducing acute phase reactants in liver, inducing fever and stimulating T cell proliferation. IL-6 activities include, for example, stimulating proliferation of mature B cells and inducing their final maturation into antibody-producing plasma cells, inducing IL-2 receptor

15 expression, inducing acute phase reactants in liver, and co-stimulation of thymocytes *in vitro*. A regulatory effect of IL-6 is inhibition of TNF α production, providing negative feedback for limiting the acute inflammatory response (Feghali, C.A. and Wright, T.M.,

20 Frontiers in Bioscience, 2, d12-26 (1997) provides a summary of cytokine activities).

The invention provides a method of inducing an immune response in an individual having a pathological condition. The method involves administering to said

25 individual an immunogenic amount of an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof.

As described above, an immunomodulatory

30 flagellin peptide can be used to beneficially boost a

general immune response in an individual having a pathological condition by stimulating an innate immune response. An increased immune response can ameliorate a pathological condition as well as prevent a pathological condition in a healthy individual, or individual not having a pathological condition. Therefore, an immunomodulatory flagellin peptide can be administered prophylactically to an individual not having a pathological condition, if desired.

10 The invention provides another method of modulating an immune response in an individual having a pathological condition. The method involves administering to the individual a combination of an immunogenic amount of an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof, and another immunomodulatory molecule.

20 As described above, a combination of an immunomodulatory flagellin peptide with another immunomodulatory molecule can be used to advantageously induce or modulate an immune response. An immune response can be induced by combining an immunomodulatory flagellin peptide with another immunomodulatory molecule that induces an immune response in a general manner, such as an adjuvant, or can be combined with an immunomodulatory molecule that induces a particular alteration in an immune cell activity. Such immunomodulatory molecules are described herein.

Modulating an immune response is useful for promoting a more effective or more normal immune response in an individual having a pathological condition. As described above, alterations in normal cytokine levels
5 are associated with various pathological conditions. An immunomodulatory flagellin peptide or combination with another immunomodulatory molecule can be used to modulate cytokine levels in an individual by inducing the production of immunomodulatory molecules, such as
10 cytokines including TNF α , IL-1, and IL-6 through TLR5, and inducing the production of suppression of the same or different immunomodulatory molecules through the activity of the administered immunomodulatory molecule. Therefore, the immunomodulatory flagellin peptides of the
15 invention can be combined with immunomodulatory molecules that alter an immune response by stimulating or inhibiting the cellular functions of immune system cells.

A variety of immunomodulatory molecules can be used in combination with an immunomodulatory flagellin
20 peptide or modification thereof of the invention to alter an immune response in an individual. The type of alteration desired will determine the type of immunomodulatory molecule selected to be combined with an immunomodulatory flagellin peptide. For example, to
25 promote an innate immune response, a immunomodulatory flagellin peptide can be combined with another immunomodulatory molecule that promotes an innate immune response, such as a PAMP or conserved region known or suspected of inducing an innate immune response. A
30 variety of PAMPs are known to stimulate the activities of different members of the toll-like family of receptors. Such PAMPs can be combined to stimulate a particular

combination of toll-like receptors that induce a beneficial cytokine profile. For example, PAMPs can be combined to stimulate a cytokine profile that induces a TH1 or TH2 immune response.

5 Other types of immunomodulatory molecules that promote humoral or cell-mediated immune responses can be combined with a flagellin molecule of the invention. For example, cytokines can be administered to alter the balance of TH1 and TH2 immune responses. Those skilled
10 in the art will know how to determine the appropriate cytokines useful for obtaining a beneficial alteration in immune response for a particular pathological condition.

 Immunomodulatory molecules that target antigens and cells displaying antigens for destruction can be
15 combined with a flagellin molecule of the invention. For example, the effectiveness of monoclonal antibodies and ADCC targeting molecules that recognize a particular antigen on an unwanted cell, such as a pathologically aberrant cell can be increased when administered with a
20 flagellin molecule of the invention. Immunomodulatory molecules that stimulate or suppress cellular activities such as proliferation, migration, activation, interaction and differentiation can be combined with a flagellin molecule of the invention. For example, IL-2 can be used
25 to stimulate proliferation of immune system cells, certain interferons can be used to interfere with the rapid growth of cancer cells or to interfere with angiogenesis, and granulocyte-colony stimulating factor can be used to increase production of certain types of
30 immune system cells and blood cells. A variety of immunostimulating and immunosuppressing molecules and

modalities are well known in the art and can be used in combination with a flagellin polypeptide, peptide or modification thereof, of the invention. A flagellin molecule of the invention increases the beneficial effect
5 of an immunomodulatory molecule by inducing TLR5-mediated production of immunomodulatory molecules that function in concert with a selected immunomodulatory molecule to produce a desired cytokine profile or cellular activity, or prime the adaptive immune response to respond to the
10 selected immunomodulatory molecule.

The methods of the invention for using immunomodulatory flagellin peptides to induce an immune response are also applicable to a flagellin polypeptide, or a modification thereof. Accordingly, the invention
15 provides a method of inducing an immune response in an individual, including a human, having a pathological condition. The method involves administering to the individual an immunogenic amount of an immunomodulatory flagellin polypeptide, or modification thereof, when the
20 flagellin polypeptide induces an immune response.

An immunomodulatory flagellin peptide of the invention binds to TLR5 and stimulates a TLR5 activity. The ability of an immunomodulatory flagellin peptide or
25 modification thereof to bind to TLR5 or stimulate a TLR5 activity can be determined using methods known in the art. Methods of determining specific binding interactions of flagellin peptides and modifications thereof with TLR5 can be determined using well known
30 methods in the art such as methods of trapping ligand-receptor complexes using chemical cross-linking, and competitive inhibition of reagents specific for TLR5 such

as specific flagellin peptides or modifications,
antibodies or other TLR-5 specific reagents.

Methods of determining TLR5 functional
activities in response to an immunomodulatory flagellin
5 peptide or modification thereof include methods described
herein, in Examples I through IV, as well as methods
known in the art. A variety of methods well known in the
art can be used for determining transcription factor
activities. For example, fos, jun, and NF- κ B activation
10 in response to TLR5 binding to a flagellin molecule can
be detected by electrophoretic mobility shift assays well
known in the art that detect NF- κ B binding to specific
polynucleic acid sequences, and promoter-reporter nucleic
acid constructs such that, for example, β -lactamase,
15 luciferase, green fluorescent protein or β -galactosidase
will be expressed in response to contacting a TLR5 with a
flagellin polypeptide, peptide or equivalent thereof.
For example, a luciferase reporter plasmid in which
luciferase protein expression is driven by one or more
20 NF- κ B binding sites can be transfected into a cell, as
described in Examples I-IV. Activation of NF- κ B results
in activation of luciferase reporter expression,
resulting in production of luciferase enzyme able to
catalyze the generation of a molecule that can be
25 detected by colorimetric, fluorescence,
chemilluminescence or radiometric assay.

An amount or activity of a polypeptide,
including a cytokine such as TNF α , IL-1 or IL-6, can be a
read-out for activation of a TLR5 in response to binding
30 an immunomodulatory flagellin peptide or modification

thereof. A variety of methods well known in the art can be used to measure cytokine amounts, such as, for example, flow cytometry methods, immunoassays such as ELISA and RIA, and cytokine RNA protection assays.

- 5 Commercially available cytokine assay kits, such as ELISA assay formats, can be conveniently used to determine the amount of a variety of cytokines in a sample. Those skilled in the art will determine the particular cytokines to be measured when assessing an immune
10 response in a cell or animal. For example, to determine whether a particular response is characterized as a TH1 or TH2 immune response, those skilled in the art will be able to select appropriate cytokines within the TH1 and TH2 categories, which are well known in the art.

- 15 A sample used for determining a TLR5-mediated response or immune response can include, for example, a fluid or tissue obtained from an animal, a cell obtained from an animal fluid or tissue, cultured cells including *in vitro* and *ex vivo* cultured cells, and lysates or
20 fractions thereof and cultured cells that express TLR5.

- An immune response in an animal is determined by the collective responses of the cells of the immune system. An immune response can be detected by observing various indicators of immune response in an animal. Such
25 indicators include, for example, visible signs of inflammation of tissues, such as swelling, production of antibodies, such as levels of IgA, IgG and IgM in blood and levels of IgA in saliva, alterations in immune cell numbers, such as increased or decreased proliferation of
30 particular immune cells, and in immune cell activities, such as production of immunomodulatory molecules and

second messenger molecules. For example, an immune response to a particular antigen can be observed in a animal using methods well known in the art such as delayed hypersensitivity skin tests. An immune response
5 can be determined by the presence of antibodies cross reactive with an antigen, such as by ELISA and Western blotting, lymphocyte activation tests employing mitogen or antigen stimulation, mixed lymphocyte culture tests, assays for human T and B lymphocytes, flow cytometry and
10 cell sorting to characterize populations of immune system cells obtained from an individual, soluble antigen uptake by macrophages, and tests of neutrophil functions (Stites et al. Basic and Clinical Immunology, 4th edition, Lange Medical Publications, Los Altos, CA (1982)). An immune
15 response can also be assessed by examining amounts or activities of immune system mediators, such as cytokines and chemokines, in cells collected from fluids or tissues of animals. A variety of methods are well known in the art for qualitative and quantitative measurement of
20 cytokine amount and bioassay of cytokine activity.

The methods of the invention for inducing an immune response can be used to treat any animal species having an immune response upon treatment with flagellin
25 polypeptide, peptide, or modification thereof, and for which a stimulation of an immune response is desired. Such animals include avian species such as chicken, and mammalian species such as rodent, canine, feline, bovine, porcine and human subjects. Methods for using adjuvants
30 with vaccines and vaccinating animals are well known in the art and are routinely used in laboratory animals. Those skilled in the art will be able to determine if a

particular animal species has a flagellin-stimulated TLR5-mediated innate immune response.

A vaccine to be used in the methods of the invention for inducing an immune response can be
5 administered as a solution or suspension together with a pharmaceutically acceptable medium. Such a pharmaceutically acceptable medium can be, for example, water, phosphate buffered saline, normal saline or other physiologically buffered saline, or other solvent or
10 vehicle such as glycol, glycerol, and oil such as olive oil or an injectable organic ester. A pharmaceutically acceptable medium can also contain liposomes or micelles, and can contain immunostimulating complexes prepared by mixing polypeptide or peptide antigens with detergent and
15 a glycoside, such as Quil A. Further methods for preparing and administering an immunomodulatory flagellin polypeptide or peptide, or modification in a pharmaceutically acceptable medium are presented below, in reference to compounds that induce a TLR-mediated
20 response.

The immunomodulatory flagellin polypeptides, peptides and modifications thereof used in the methods of the invention can be administered by a variety of routes to stimulate an immune response. For example, these
25 immunomodulatory molecules can be delivered intranasally, subcutaneously, intradermally, intralymphatically, intramuscularly, intratumorally, orally, intravesically, intraperitoneally and intracerebrally. Oral administration is convenient and relatively safe. Oral
30 vaccination protocols can be useful for inducing the state of immunological tolerance which normally occurs in

response to most soluble antigens and the proteolytic degradation of antigen preparations in the digestive tract. Nasal delivery routes may be useful for inducing both mucosal and systemic immune responses. A variety of
5 devices are under development for convenient and effective delivery of formulations to the nasal cavity and pulmonary tissues. Those skilled in the art will know how to select appropriate delivery routes for particular formulations of flagellin polypeptides,
10 peptides and modifications thereof.

The invention provides a screening composition consisting of a flagellin peptide of claim 1 and a TLR5. The composition is useful for identifying agonists, antagonists and ligands for TLR5. The characteristics of
15 a flagellin peptide of claim 1 and preparation of a flagellin peptide are described herein. Similarly, the characteristics of a TLR5 polypeptide and modifications thereof that have a TLR5 activity, and methods for preparing a TLR5 polypeptide to be used in the methods of
20 the invention are described herein. Chimeric TLR5s, such as the CD4-TLR5 described herein in Example I, are included in the screening compositions of the invention.

The screening composition of the invention includes, for example, cells, cell extracts and
25 artificial signaling systems that contain a TLR5 polypeptide or modification thereof. The cell compositions of the invention include any cell in which TLR5 can couple to a signal transduction pathway to produce a detectable signal in response to an agonist,
30 such as flagellin or a flagellin peptide. Such cells include insect cells such as *Drosophila* cells, yeast

cells such as *S. cerevisiae*, prokaryotic cells such as *E. coli*, amphibian cells such as *Xenopus* oocytes, and vertebrate cells such as mammalian primary cells, such as macrophages. Primary cells such as macrophages and other lymphocytes can be conveniently isolated from blood using methods well known in the art. Cells obtained from transgenic animals, such as transgenic mice that have been engineered by known methods of express recombinant TLR5 or TLR5 signal transduction components are also included in the screening compositions of the invention. Cell lines prepared from any of theses cell types, such as S2, CHO, NIH-3T3, 293 and HeLa cells are also included in a screening composition of the invention.

The screening compositions of the invention can include crude or partially purified lysates or extracts of the cell compositions of the invention, and reconstituted signaling systems. Artificial signaling systems include, for example, natural or artificial lipid bilayers, such as a liposome or micelle, which promote an active conformation of a TLR5. The compositions can further contain cellular fractions or isolated components necessary for producing and detecting the desired predetermined signal.

The invention provides a method of screening for a TLR5 ligand, agonist or antagonist. The method involves, (a) contacting a TLR5 with a candidate compound in the presence of a flagellin polypeptide or immunomodulatory flagellin peptide under conditions wherein binding of the flagellin polypeptide or immunomodulatory flagellin peptide to the TLR5 produces a predetermined signal; (b) determining the production of

the predetermined signal in the presence of the candidate compound; and (c) comparing the predetermined signal in the presence of the candidate compound with a predetermined signal in the absence of the candidate compound, wherein a difference between the predetermined signals in the presence and absence of the candidate compound indicates that the compound is a TLR5 ligand, agonist or antagonist.

TLR5 can produce a variety of predetermined signals useful in the methods of the invention for identifying a TLR5 ligand, agonist or antagonist. TLR5 has an extracellular domain that participates in ligand recognition and intracellular domain that contain a conserved region called the Toll/IL-1R homology (TIR) domain that, upon activation, recruits an adaptor protein, MyD88. Through an amino terminal death domain, MyD88 recruits the serine kinase IRAK to propagate a pro-inflammatory signal through binding to TRAF6, which then binds to other molecules that participate in the TLR5 signaling cascade. Immunomodulatory flagellin peptides and modifications binding to TLR5 induces signal transduction events which result in, for example, stimulating NF- κ B activity and inducing production of gene products of NF- κ B-regulated genes, such as TNF α , IL-1 and IL-6, as well as stimulating AP-1 transcription factors fos and jun. Therefore, a predetermined signal can include a signal produced by an immunomodulatory flagellin polypeptide or peptide or modification binding to TLR5, a signal produced by a TLR5 intracellular signal transduction even, such as kinase or phosphatase activity or protein-protein interactions, by activation of fos,

jun or NF- κ B, and by an amount or activity of a fos-, jun- or NF- κ B-regulated gene or gene product, such as TNF α , IL-1 and IL-6.

5 A variety of low- and high-throughput assays suitable for detecting selective binding interactions between a receptor and a ligand are known in the art. Both direct and competitive assays can be performed, including, for example, fluorescence correlation
10 spectroscopy (FCS) and scintillation proximity assays (SAP) reviewed in Major, J. Receptor and Signal Transduction Res. 15:595-607 (1995); and in Sterrer et al., J. Receptor and Signal Transduction Res. 17:511-520 (1997)). Other assays for detecting binding interactions
15 include, for example, ELISA assays, FACS analysis, and affinity separation methods. Such assays can involve labeling a TLR5 ligand, such as flagellin or a flagellin peptide, with a detectable moiety such as a radiolabel, fluorochrome, ferromagnetic substance, or luminescent
20 substance. A detectably labeled flagellin polypeptide or peptide can be prepared using methods well known in the art. Receptor binding assays, including high-throughput automated binding assays, and methods of determining binding affinity from such assays, are well known in the
25 art, and any suitable direct or competitive binding assay can be used. Exemplary high-throughput receptor binding assays are described, for example, in Mellentin-Micelotti et al., Anal. Biochem. 272:P182-190 (1999); Zuck et al., Proc. Natl. Acad. Sci. USA 96:11122-11127 (1999); and
30 Zhang et al., Anal. Biochem. 268:134-142 (1999).

A variety of methods well known in the art can be used to detect activation of transcription factors, such as NF- κ B, in low- or high-throughput formats. The methods described herein and in the Examples can be
5 adapted to formats suitable for candidate compound screening.

A variety of low- and high-throughput assays suitable for detecting amounts and activities of polypeptides such as cytokines are known in the art.
10 Methods for detecting polypeptides, include, for example, flow cytometric measurements as described herein, immunodetection methods such as radioimmune assay (RIA), ELISA, immunoprecipitation and Western blotting. Assay of the activity of a cytokine include function bioassays
15 and detection of amounts of polypeptides regulated by a particular cytokine. Those skilled in the art can determine an appropriate method for detecting an activity of a particular cytokine.

Suitable conditions under which TLR5 produces a
20 predetermined signal in response to a flagellin polypeptide, peptide or modification can be determined by those skilled in the art, and will depend on the particular predetermined signal selected. Exemplary conditions for determining the production of a
25 predetermined signal are provided herein in Examples I-IV. Any known or predicted TLR5-mediated cellular event, such as elicitation of second messengers, induction of gene expression or altered cellular proliferation, differentiation or viability can be a predetermined
30 signal that is an indication of activation of signal transduction through TLR5.

Assays for detecting a predetermined signal produced by binding of flagellin or flagellin peptide to TLR5 can be performed, for example, with whole cells that express TLR5, membrane fractions, or artificial systems, as described herein, or with isolated TLR5 polypeptide, either in solution, in an artificial membrane, or bound to a solid support.

A method of identifying TLR5 agonists and antagonists can be performed either in the presence of a predetermined concentration of a known TLR5 agonist, such as flagellin, flagellin peptide, or modifications thereof, or in the absence of agonist. The agonist can be added either prior to, simultaneously with, or after, addition of the test compound. When present, the agonist concentration is preferably within 10-fold of its EC50 under the assay conditions to allow the identification of a compound that competes with a known agonist for signaling through TLR5, or indirectly augments signaling through the receptor. Likewise, a compound that reduces binding between a known agonist and its receptor, or indirectly decreases signaling through the receptor, can also be identified.

The method of screening to identify a ligand, agonist or antagonist of TLR5 involve testing a candidate compound. A candidate compound can be any substance, molecule, compound, mixture of molecules or compounds, or any other composition. The candidate compounds can be small molecules or macromolecules, such as biological polymers, including proteins, polysaccharides and nucleic acids. Sources of candidate compounds which can be

screened for a ligand, agonist or antagonist of TLR5 include, for example, libraries of small molecules, peptides and polypeptides.

Additionally, candidate compounds can be
5 preselected based on a variety of criteria. For example, suitable candidate compounds can be selected as having known ligand, agonist or antagonist activity. Alternatively, candidate compounds can be selected randomly. Candidate compounds can be administered to the
10 reaction system at a single concentration or, alternatively, at a range of concentrations to determine, for example, an EC50 or IC50 of a candidate compound.

The method of screening for TLR5 ligands, agonists or antagonists can involve groups or libraries
15 of compounds. Methods for preparing large libraries of compounds, including simple or complex organic molecules, carbohydrates, peptides, peptidomimetics, polypeptides, nucleic acids, antibodies, and the like, are well known in the art. Libraries containing large numbers of
20 natural and synthetic compounds can be obtained from commercial sources.

The number of different candidate compounds to examine using the methods of the invention will depend on the application of the method. It is generally
25 understood that the larger the number of candidate compounds, the greater the likelihood of identifying a compound having the desired activity in a screening assay. Large numbers of compounds can be processed in a high-throughput automated format.

The TLR5 agonists, antagonists and ligands identified using the methods and compositions described herein, are potential therapeutic compounds that can be administered to an individual, such as a human or other mammal, in an effective amount to increase or decrease signaling through TLR5, for example, to alter an immune response or treat a TLR5-associated condition. Such compounds can be used analogously to immunomodulatory compounds useful for augmenting and altering an immune response, as described above. For example, a compound can be used to induce a general immune response and to induce a specific immune response in the presence of an antigen and to alter the level of a particular cytokine in an individual having a pathological condition.

15

The TLR5 agonists and antagonists, immunomodulatory flagellin peptides, polypeptides and modifications thereof, are useful for ameliorating, or reducing the severity of a pathological condition. Reduction in severity includes, for example, an arrest or decrease in clinical symptoms, physiological indicators, biochemical markers or metabolic indicators of disease. Those skilled in the art will know, or will be able to determine the appropriate clinical symptoms, physiological indicators, biochemical markers or metabolic indicators to observe for a particular pathological condition. To prevent a disease means to preclude the occurrence of a disease or restoring a diseased individual to their state of health prior to disease.

30

In addition to applications described herein for agonists and antagonists, a TLR5 ligand can be used,

for example, to specifically target a diagnostic moiety to cells and tissues that express TLR5, such as monocytes, immature dendritic cells, epithelial cells, and other cells involved in an immune response. Thus, a
5 TLR5 ligand can be labeled with a detectable moiety, such as a radiolabel, fluorochrome, ferromagnetic substance, or luminescent substance, and used to detect normal or abnormal expression of TLR5 polypeptide in an isolated sample or *in vivo* diagnostic imaging procedures.

10 A heterologous amino acid sequence can be advantageously used to provide a tag for detection or purification or to impart an activity to a reference polypeptide or peptide, such as an enzyme activity, a biological activity, an immunological activity or
15 stability. An immunomodulatory flagellin peptide, polypeptide or modification thereof, or TLR5 polypeptide can contain a heterologous amino acid sequence, or amino acid sequence not present in the native amino acid sequence of a reference polypeptide or peptide and not
20 represented by a modification of a reference polypeptide or peptide. A heterologous amino acid sequence can be of any size in relation to the reference amino acid sequence. A TLR5 polypeptide containing the heterologous sequence of CD4 is a specific example of such a
25 modification and is described further in Example I. The described CD4-TLR5 chimera is identified by the amino acid sequence of SEQ ID NO:8, encoded by the nucleic acid sequence of SEQ ID NO:7. A chimeric TLR5 can be prepared using cloning methods well known in the art. For
30 example, a chimeric polypeptide can be produced by amplifying by PCR a nucleotide sequence encoding a portion of a selected polypeptide using sequence specific

primers. Primers useful for amplifying a TLR5 include, for example, huTLR5-A6:

TTAAAGTGGTACCAGTTCTCCCTTTTCATTGTATGCACT and

huTLR5DNS: CGGGATCCCGTTAGGAGATGGTTGCTACAGTTTGC. A

5 portion of a TLR5 nucleotide sequence, such as a sequence amplified using such primers can be fused to a nucleotide sequence encoding a heterologous amino acid sequence. A variety of methods for generating nucleic acid sequences encoding chimeric polypeptides are well known to those
10 skilled in the art.

The polypeptides and peptides described herein, including immunomodulatory flagellin peptides, flagellin polypeptide, TLR5 polypeptides and fragments thereof can
15 be prepared using a variety of protein expression systems well known in the art, including prokaryotic and eukaryotic expression systems. Prokaryotic expression systems are advantageous due to their ease in manipulation, low complexity growth media, low cost of
20 growth media, rapid growth rates and relatively high yields. Well known prokaryotic expression systems include, for example, *E. coli* bacterial expression systems based on bacteriophage T7 RNA polymerase, the *trc* promoter, the *araB* promoter and bacillus expression.

25 Eukaryotic expression systems are advantageous because expressed polypeptides can contain eukaryotic post-translational modifications such as O-linked glycosylation, phosphorylation and acetylation and can have improved protein folding. Well known eukaryotic
30 expression systems include, for example, expression in yeast, such as *Pichia pastoris* and *Pichia methanolica*, expression in insect systems such as the *Drosophila* S2 system and baculovirus expression systems and expression

in mammalian cells using adenoviral vectors and cytomegalovirus promoter-containing vectors.

An immunomodulatory flagellin peptide, polypeptide, TLR5 or fragments thereof can be purified
5 using a variety of methods of protein purification well known in the art. Biochemical purification can include, for example, steps such as solubilization of the polypeptide or peptide-expressing cell, isolation of the desired subcellular fractions, chromatography, such as
10 ion exchange, size, or affinity-based chromatographies, electrophoresis, and immunoaffinity procedures. Other well-known methods are described in Deutscher et al., Guide to Protein Purification: Methods in Enzymology Vol. 182, (Academic Press, (1990)). An exemplary method for
15 purifying a flagellin peptide is provided in Example III. The methods and conditions for biochemical purification of a polypeptide of the invention can be chosen by those skilled in the art, and the purification monitored, for example, by staining SDS-PAGE gels containing protein
20 samples, by immunodetection methods such as Western blotting and ELISA, and by functional assay of immunogenic activity of flagellin or a TLR5 activity of TLR5.

An immunomodulatory flagellin peptide,
25 polypeptide, TLR5 or fragments thereof can be modified, for example, to increase polypeptide stability, alter an activity, facilitate detection or purification, or render the enzyme better suited for a particular application, such as by altering substrate specificity. Computer
30 programs known in the art can be used to determine which amino acid residues of a immunomodulatory flagellin

peptide, flagellin polypeptide or TLR5 can be modified as described above without abolishing a corresponding activity (see, for example, Eroshkin et al., Comput. Appl. Biosci. 9:491-497 (1993)). In addition, structural and sequence information can be used to determine the amino acid residues important for activity. For example, a comparisons of flagellin amino acid sequences, such as that shown in Figure 7 can provide guidance in determining amino acid residues that can be altered without abolishing flagellin or flagellin peptide activity by indicating amino acid residues that are conserved across species. Conserved regions of flagellin are well known in the art and have been described, for example, in Mimori-Kiyosue, et al., J. Mol. Biol. 270:222-237, (1997). A crystal structure of flagellin can also provide guidance for making flagellin modifications (Samatey et al. Nature, 410:331-337 (2001)). Similarly, amino acid sequence comparisons between the disclosed murine TLR5, TLR5s of other species, and other toll-like receptor family members can provide guidance for determining amino acid residues important for activity.

An isolated TLR5 is a TLR5 removed from one or more components with which it is naturally associated. Therefore, an isolated TLR5 can be a cell lysate, cell fraction, such as a membrane fraction, or a purified TLR5 polypeptide. An isolated TLR5 can include a liposome or other compound or matrix that stabilizes or promotes an active conformation of the receptor.

For treating or reducing the severity of a pathological condition a TLR5 agonist or antagonist,

immunomodulatory flagellin peptide, polypeptide or modification thereof, including a vaccine, can be formulated and administered in a manner and in an amount appropriate for the condition to be treated; the weight, gender, age and health of the individual; the biochemical nature, bioactivity, bioavailability and side effects of the particular compound; and in a manner compatible with concurrent treatment regimens. An appropriate amount and formulation for a particular therapeutic application in humans can be extrapolated based on the activity of the compound in recognized animal models of the particular disorder.

Animal models of aberrantly proliferative diseases can be used to assess a formulation of compound, including a vaccine or adjuvant containing an immunomodulatory flagellin peptide, polypeptide or modification thereof, for an amount sufficient to induce an immune response or ameliorate disease symptoms. Animal models of such pathological conditions well known in the art which are reliable predictors of treatments in human individuals for include, for example, animal models for tumor growth and metastasis, infectious diseases and autoimmune disease.

There are numerous animal tumor models predictive of therapeutic treatment which are well known in the art. These models generally include the inoculation or implantation of a laboratory animal with heterologous tumor cells followed by simultaneous or subsequent administration of a therapeutic treatment. The efficacy of the treatment is determined by measuring the extent of tumor growth or metastasis. Measurement of

clinical or physiological indicators can alternatively or additionally be assessed as an indicator of treatment efficacy. Exemplary animal tumor models can be found described in, for example, Brugge et al., Origins of
5 Human Cancer, Cold Spring Harbor Laboratory Press, Plain View, New York, (1991).

Similarly, animal models predictive for infectious disease also follow a similar approach. Briefly, laboratory animals are inoculated with an
10 infectious agent and the progression of the infection is monitored by, for example, clinical symptoms, growth culture of the agent from an infected tissue sample or biopsy in the presence or absence of the therapeutic
15 treatment. The reduction in severity of the diagnostic indicator is indicative of the efficacy of the treatment. A variety of animal models for infectious diseases are well known to those skilled in the art.

One animal model predictive for autoimmune diseases is Experimental allergic encephalomyelitis
20 (EAE), also called experimental autoimmune encephalomyelitis. Although originally characterized as a model for neurological autoimmune disease such as human multiple sclerosis, the use of this model to predict
25 treatments of other autoimmune diseases has been widely accepted. EAE is induced in susceptible animals by active immunization with myelin basic protein (MBP) or by passive transfer of MBP-specific T helper lymphocytes. Progression of the disease is characterized by chronic relapsing paralysis and central nervous system
30 demyelination, which can be monitored by observation or by immunological determinants such as delayed-type

hypersensitivity (DTH; a measure of cell mediated immunity) response to the immunogen. Efficacy of a therapeutic treatment is compared to progression of the disease in the absence of treatment. A reduction in
5 severity of EAE symptoms or immunological determinants in treated animals is indicative of the efficacy of the therapeutic treatment. For a review of autoimmune disease models see, for example, Urban et al., *Cell*, 54:577-592 (1988); Brostoff et al., *Immunol. Ser.* 59:203-
10 218 (1993) and U.S. Patent Nos. 5,614,192 and 5,612,035.

A growing number of human diseases have been classified as autoimmune and include, for example, rheumatoid arthritis, myasthenia gravis, multiple sclerosis, psoriasis, systemic lupus erythmatosis,
15 autoimmune thyroiditis, Graves' disease, inflammatory bowel disease, autoimmune uveoretinitis, polymyositis and diabetes. Animal models for many of these have been developed and can be employed analogously as the EAE model described above predictive assessment of
20 therapeutic treatments using the compounds, vaccines and adjuvants in the methods of the invention.

Other reliable and predictive animal models are well known in the art and similarly can be used to assess a compound formulation, including vaccine and adjuvant
25 formulations containing an immunomodulatory flagellin peptide, polypeptide or modification thereof.

The total amount of a compound including an immunomodulatory flagellin peptide, polypeptide or modification thereof, that modulates a TLR5-mediated
30 immune response can be administered as a single dose or

by infusion over a relatively short period of time, or can be administered in multiple doses administered over a more prolonged period of time. Additionally, a compound can be administered in a slow-release matrix, which can
5 be implanted for systemic delivery at or near the site of the target tissue.

A compound that modulates a TLR5-mediated immune response can be administered to an individual using a variety of methods known in the art including,
10 for example, intravenously, intramuscularly, subcutaneously, intraorbitally, intracapsularly, intraperitoneally, intracisternally, intra-articularly, intracerebrally, orally, intravaginally, rectally, topically, intranasally, or transdermally.

15 A compound that modulates a TLR5-mediated immune response can be administered to a subject as a pharmaceutical composition comprising the compound and a pharmaceutically acceptable carrier. The choice of pharmaceutically acceptable carrier depends on the route
20 of administration of the compound and on its particular physical and chemical characteristics. Pharmaceutically acceptable carriers are well known in the art and include sterile aqueous solvents such as physiologically buffered saline, and other solvents or vehicles such as glycols,
25 glycerol, oils such as olive oil and injectable organic esters. A pharmaceutically acceptable carrier can further contain physiologically acceptable compounds that stabilize the compound, increase its solubility, or increase its absorption. Such physiologically acceptable
30 compounds include carbohydrates such as glucose, sucrose or dextrans; antioxidants, such as ascorbic acid or

glutathione; chelating agents; and low molecular weight proteins. As described above in reference to vaccines, such routes of administration are also applicable to administration of an immunomodulatory flagellin peptide,
5 polypeptide or modification thereof.

In addition, a formulation of a compound that modulates a TLR5-mediated immune response can be incorporated into biodegradable polymers allowing for sustained release of the compound, the polymers being
10 implanted in the vicinity of where drug delivery is desired, for example, at the site of a tumor or implanted so that the compound is released systemically over time. Osmotic minipumps also can be used to provide controlled delivery of specific concentrations of a compound through
15 cannulae to the site of interest, such as directly into a tumor growth or other site of a pathology involving a perturbation state. The biodegradable polymers and their use are described, for example, in detail in Brem et al., J. Neurosurg. 74:441-446 (1991). These methods, in
20 addition to those described above in reference to vaccines, are applicable to administering an immunomodulatory flagellin peptide, polypeptide or modification thereof to induce an immune response.

The methods of treating a pathological
25 condition additionally can be practiced in conjunction with other therapies. For example, for treating cancer, the methods of the invention can be practiced prior to, during, or subsequent to conventional cancer treatments such as surgery, chemotherapy, including administration
30 of cytokines and growth factors, radiation or other methods known in the art. Similarly, for treating

pathological conditions which include infectious disease, the methods of the invention can be practiced prior to, during, or subsequent to conventional treatments, such as antibiotic administration, against infectious agents or
5 other methods known in the art. Treatment of pathological conditions of autoimmune disorders also can be accomplished by combining the methods of the invention for inducing an immune response with conventional treatments for the particular autoimmune diseases.
10 Conventional treatments include, for example, chemotherapy, steroid therapy, insulin and other growth factor and cytokine therapy, passive immunity and inhibitors of T cell receptor binding. The methods of the invention can be administered in conjunction with
15 these or other methods known in the art and at various times prior, during or subsequent to initiation of conventional treatments. For a description of treatments for pathological conditions characterized by aberrant cell growth see, for example, The Merck Manual, Sixteenth
20 Ed, (Berkow, R., Editor) Rahway, N.J., 1992.

As described above, administration of a compound, immunomodulatory flagellin peptide, flagellin polypeptide or modification thereof can be, for example, simultaneous with or delivered in alternative
25 administrations with the conventional therapy, including multiple administrations. Simultaneous administration can be, for example, together in the same formulation or in different formulations delivered at about the same time or immediately in sequence. Alternating
30 administrations can be, for example, delivering an immunomodulatory flagellin peptide or polypeptide formulation and a conventional therapeutic treatment in

temporally separate administrations. As described previously, the temporally separate administrations of a compound, immunomodulatory flagellin peptide, polypeptide or modification thereof, and conventional therapy can
5 similarly use different modes of delivery and routes.

The invention provides a method of using a signal produced in response to flagellin binding to TLR5 to detect bacterial contamination in a sample. The method can be used to detect picogram amounts of
10 flagellin in a sample.

Food-borne diseases resulting from the presence of harmful bacteria account for 325,000 hospitalizations and 5,000 deaths each year in the United States (National Institutes of Health, Foodborne Diseases NIAID Fact
15 Sheet). The U.S. Centers for Disease Control and Prevention (CDC) estimates that 1.4 million people in the United States are infected each year with *Salmonella*. Other bacterial pathogens that cause pathological conditions characterized by symptoms ranging from
20 intestinal discomfort to severe dehydration, bloody diarrhea and even death, include enterohemorrhagic *E. coli*, such as strains designated O157:H7 and O26:H11, *Campylobacter* strains such as *C. jejuni*, and *Shigella* strains such as *S. flexneri*.

25 All of these bacterial strains are flagellated, and therefore express flagellin polypeptides. For example, the amino acid sequences of flagellins from *Salmonella*, *E. coli*, *Campylobacter*, *Shigella* strains are shown in Figure 7. The methods of the invention for
30 detecting flagellin polypeptides contained in samples

suspected of bacterial contamination can be applied to quality assurance protocols for preparation of foods and numerous other applications.

The invention also provides a bioassay for
5 detecting bacterial contamination in a sample. The method involves, (a) contacting the sample with a TLR5 under conditions wherein binding of a flagellin polypeptide or fragment thereof in the sample to the TLR5 produces a predetermined signal, (b) determining the
10 production of the predetermined signal in the presence and absence of the sample, and (c) comparing the predetermined signal in the presence of the sample with a predetermined signal in the absence of the sample, wherein a difference between the predetermined signals in
15 the presence and absence of the sample indicates that the sample contains flagellin.

The methods of the invention for detecting bacterial contamination are based on the finding disclosed herein that flagellin is a ligand for TLR5.
20 Therefore, a flagellin molecule in a sample can bind to a TLR5 and elicit the production of a predetermined signal. A predetermined signal produced by TLR5 in a particular assay system is compared in the presence and absence of a sample known or suspected of containing a bacterial
25 contaminant. A sample known to be free of flagellin can be used as a negative control, while a sample containing a known concentration of flagellin, flagella or bacteria having flagella can be used as a positive control.

A sample to be tested for the presence of
30 flagellin can be any material that is suspected of being

contaminated with a gram-positive or gram-negative flagellated bacterium. For example, the method for determining the presence of flagellin can be performed using a sample of a biological fluid, cell, tissue, organ
5 or portion thereof, such as a sample of a tissue to be used for preparing a product, a product for human or animal consumption, such as a food or pharmaceutical preparation, and a product for external application or administration by any route to an animal.

10 A variety of predetermined signals produced by a TLR5, as discussed above and in the Examples herein, can be used to detect the binding and activation of a TLR5 by a flagellin molecule present in a sample. A variety of methods known in the art, including those
15 described herein can be used to detect a predetermined signal produced by a TLR5.

It is understood that modifications which do not substantially affect the activity of the various
20 embodiments of this invention are also included within the definition of the invention provided herein. Accordingly, the following examples are intended to illustrate but not limit the present invention.

EXAMPLE I

Constitutively Active TLR5 Activates NF- κ B
and TNF α Production

This example shows activation of NF- κ B and TNF α production in CHO cells in response to constitutively active TLR5.

To determine if TLR5 activates NF- κ B and TNF α production, the activity of a constitutively active form of TLR5 was examined in CHO cells. Constitutively active forms of TLR4 and TLR5 were generated by fusing the extracellular domain of CD4 to the transmembrane and TIR domain of TLR4 or TLR5 (Medzhitov, R. et al. Nature 388, 394-7 (1997); Ozinsky, A. et al., Proc. Natl. Acad. Sci. 97, 13766-13881 (2000)). CD4-TLR5 was constructed by fusing the murine CD4 extracellular domain (amino acids 1-391) to the putative transmembrane and cytoplasmic domains of human TLR5 (amino acids 639-859) and cloning into pEF6-TOPO (pEF6-mCD4-hTLR5). These chimeras, referred to as CD4-TLR4 and CD4-TLR5 were expressed in CHO cells.

For determining NF- κ B activity in response to TLR5, CHO cells were transiently transfected with expression vectors for CD4-TLR4, CD4-TLR5, or empty expression vector (control) together with an NF- κ B luciferase reporter. NF- κ B-induced luciferase activity was measured. CHO cells (CHO-K1) were obtained from ATCC (no. CRL-9618) and grown in Ham's F-12 medium supplemented with 10% FBS, L-glutamine, penicillin, and

streptomycin. CHO cells were transfected by electroporation as described previously (Underhill, D. M. et al., Nature, 401,811-5 (1999)), with 1 µg of the indicated TLR expression vector, 1 µg of ELAM-firefly luciferase, 0.1 µg of TK-renilla luciferase (Promega). Cells were plated on 96-well plates at 100,000 cells/well, and incubated overnight at 37°C, 5% CO₂. Firefly and renilla luciferase activities were measured using the Dual Luciferase Assay System (Promega, Madison, WI). Luciferase activity is expressed as a ratio of NF-κB-dependent ELAM-firefly luciferase activity divided by control thymidine kinase-renilla luciferase activity (relative luciferase units).

For determining TNFα production in response to TLR5, RAW-TTIO Macrophage cells were transfected with a CD4-TLR5 expression vector, and the production of TNFα was measured by flow cytometry, as described previously (Ozinsky, A. et al. Proc. Natl. Acad. Sci. 97, 13766-13771 (2000)). Transfections were performed by electroporation using 10 µg of pEF6-mCD4-hTLR5, and 18 hours later the cells were incubated with 5 µg/ml of brefeldin A for 4 hours to accumulate intracellular pools of newly synthesized TNFα. Cells were fixed, permeabilized, stained for the expression of CD4 (anti-CD4-FITC, Pharmingen) and TNFα (anti-murine TNFα-PE, Pharmingen), and analyzed on a FACScan (Beckton-Dickenson). FACS data were analyzed with WinMDI (Joseph Trotter, Scripps Research Institute, La Jolla, CA). Cells were gated to exclude dead cells and for expression of CD4.

Figure 1 shows that expression of CD4-TLR5 induced NF- κ B activation-mediated luciferase production in CHO cells (Fig. 1a) and TNFA- α production in mouse macrophages (Fig. 1b). In Figure 1b, the dotted line indicates TNFA- α produced in cells not expressing CD4-TLR5, and the solid line indicates TNFA- α produced in cells expressing CD4-TLR5.

Thus, homo-oligomerization of the TLR5 signaling domain induces a cellular signal characterized by the induction of NF- κ B activity and production of TNF α .

EXAMPLE II

Bacterial Culture Supernatants Contain TLR5-Stimulating Activity

This Example shows that bacterial culture supernatants contain TLR5-stimulating activity.

CHO cells expressing human TLR5 and a luciferase-linked reporter were used to screen for PAMPs recognized by the receptor. PAMPs tested included LPS, lipopeptide, yeast, and extracts from *E.coli*, *Pseudomonas*, and *Listeria*. CHO cells were transiently transfected with TLR2, TLR5, or empty expression vectors together with a NF- κ B luciferase reporter. The cells were treated with 100 ng/ml LPS, 100 ng/ml lipopeptide, 10⁷ yeast particles/ml, or untreated (control), and luciferase activity was measured. The cells were treated with 67 μ g/ml of supernatant from the indicated saturated

bacterial cultures, or LB alone (control), and the luciferase activity was measured. Data are representative of 3 independent experiments.

Human TLR5 and TLR2 were generated by PCR from
5 cDNA derived from human peripheral blood mononuclear cells and cloned into pEF6-TOPO (Invitrogen) (pEF6-hTLR5 and pEF6-hTLR2). Murine TLR5 was generated by PCR using cDNA derived from RAW-TT10 cells and cloned into pEF6 (pEF6- mTLR5).

10 For luciferase assays, CHO cells were transfected by electroporation as described above, with 1 µg of the indicated TLR expression vector, 1 µg of ELAM-firefly luciferase, 0.1 µg of TK-renilla luciferase (Promega, Madison, WI). The medium was replaced with
15 medium containing the stimuli at the indicated concentration/dilution. Bacterial lipopeptide and PAM₃CSK₄, were obtained from Roche, LPS (*Salmonella minnesota* R595) was from List, and yeast particles (zymosan) were from Molecular Probes (Eugene, OR). Cells
20 were stimulated for 5 hours at 37°C, and firefly and renilla luciferase activities were measured using the Dual Luciferase Assay System (Promega).

For preparation of bacterial supernatants, bacteria were grown either in Luria broth (LB)
25 (*Escherichia coli* TOP 10 (Invitrogen), *Salmonella minnesota* (ATCC#49284), mutant *Salmonella typhimurium* (TH4778 *fliB*⁻ *fliC*⁺), TH2795 (*fliB*⁻ *fliC*⁻), (Dr. Kelly Hughes, University of Washington), or grown in trypticase soy broth (TSB) (*Listeria monocytogenes* (10403, gift of

Dr. Daniel Portnoy, UCSF), *Listeria innocua* (ATCC#33090), *Bacillus subtilis*, and *Pseudomonas aeruginosa* (Susan R. Swanzy, University of Washington)). Bacteria were grown to saturation (about 16 hours, 37°C with vigorous
5 aeration). The bacterial culture supernatants were centrifuged for 30 minutes at 2000 x g, filtered (0.2 µM), and stored at 4°C prior to use. For *flaA* transfections, *E.coli* TOP10 containing pTrcHis2-*flaA* or pTrcHis2-*flaA*rev were selected from bacterial plates and
10 grown to OD₆₀₀ of 0.6 in LB with 100 µg/ml ampicillin and 1% w/v glucose. The bacteria were centrifuged for 30 minutes at 2000 x g, and split into two LB cultures, one containing 100 µg/ml ampicillin and 1% w/v glucose (to repress *flaA*) and the other containing 100 µg/ml
15 ampicillin and 1 mM IPTG (to induce *flaA*). Samples were taken at 4 hours after induction, centrifuged 5 min at 10,000 x g, and the supernatants stored at 4°C before use.

TLR5 did not respond to any of the PAMPs known to stimulate TLR pathways, such as LPS, lipopeptide,
20 yeast cell wall, or peptidoglycan, while CHO cells transfected with TLR2 were stimulated by lipopeptide, yeast cell wall, and peptidoglycan (Fig. 2a). However, TLR5-stimulating activity was detected in culture supernatants of a variety of Gram-positive and
25 Gram-negative bacteria (Fig. 2b). The TLR5-stimulating activity of Gram-positive bacteria was not enhanced by co-expression of CD14. Interestingly, the TOP10 strain of *E. coli* had very little TLR5 activity (Fig. 2b), and was used in subsequent reconstitution experiments (see
30 below). Experiments using murine TLR5 yielded similar results.

Thus, the activity of TLR5 was stimulated by a component of bacterial culture supernatants, but not by PAMPs known to stimulate other toll like receptor family members.

5

EXAMPLE III

Purification of TLR5-Stimulating Activity from *L. monocytogenes* Culture Supernatant

This Example shows the purification of
10 TLR5-simulating activity from *L. monocytogenes* culture supernatant.

The biological activity recognized by TLR5 was determined to be TCA precipitable, phenol soluble, and sensitive to proteinase K and trypsin digestion. To
15 identify the bacterial components that stimulate TLR5, the supernatant from a saturated *L. monocytogenes* culture was concentrated, fractionated by reverse-phase chromatography, and each fraction was assessed for TLR5-stimulating activity in CHO cells (Fig. 3a).

20 For assessing the TLR-stimulating activity of FPLC fractions, CHO cells were transfected as described in Example I with the addition of 0.1 µg of pNeo/Tak (Underhill et al., Nature 401, 811-5 (1999)), and stable populations of cells expressing the indicated TLR with
25 the luciferase reporters were selected in 100 µg/ml G418. These cells were plated on 96-well plates at 100,000 cells/well and incubated overnight.

For the purification of the TLR5-stimulating activity, saturated *L. monocytogenes* culture (200 ml of TSB) was centrifuged, and the supernatant was enriched for molecules larger than 30 kDa by ultrafiltration (Ultrafree-15 filter unit with Biomax-30 membrane, Millipore). The buffer was changed to 100 mM Tris pH 7.5, and the volume was adjusted to 5 ml. The sample was loaded onto a HR5/10 reverse-phase chromatography column (AP Biotech) and run at 0.3 ml/min. Reverse-phase chromatography was performed with the indicated elution profile using the following buffers: (A) initial buffer, 0.1% TFA in water, (B) final buffer, 0.1% TFA in acetonitrile. Fractions were collected at 3-minute intervals. FPLC fractions (50 μ l) were separated on a 10% SDS-PAGE gel.

As shown in Figure 3a, CHO cells expressing an NF- κ B luciferase reporter and TLR5 were stimulated with reverse-phase FPLC fractions, and TLR5-mediated NF- κ B luciferase activity was measured. The fraction numbers correspond to 3 minute fractions of reverse-phase FPLC eluted with a non-linear gradient of buffer B as shown. Fraction number "N" is control LB growth medium and "P" is the *L. monocytogenes* culture supernatant prior to chromatography. Fractions containing background activity (1), low activity (2) and high activity (3) as indicated in Fig. 3a were analyzed by SDS-PAGE and silver stain. Silver staining was performed according to established methods. Two bands with apparent molecular masses of 30-34 kDa were clearly enriched in the fraction containing the highest level of TLR5-stimulating activity (Fig. 3b, Lane 3). Proteins eluted from regions A, B,

and C of the SDS-PAGE gel, as indicated in Fig. 3b were assayed for TLR5-mediated NF- κ B activation in CHO cells. In Fig. 3c, "Listeria" indicates *L. monocytogenes* culture supernatant. One of these bands, (Fig. 3b, band A), was
5 trypsin-treated, subject to microcapillary HPLC-tandem mass spectrometry, and identified by comparison of peptide tandem mass spectra to sequences in a non redundant protein database using the computer program, SEQUEST27 (Fig. 4a). TLR5-stimulating activity was not
10 recovered from any other section of the gel.

Thus, a TLR5-stimulating activity was purified from culture supernatants from *L. monocytogenes*.

EXAMPLE IV

Flagellin is a TLR5 Stimulus

15 This example shows that flagellin is a TLR5 stimulus purified from culture supernatants from *L. monocytogenes*.

As described above, a TLR5-stimulating activity was purified from *L. monocytogenes* culture supernatants
20 using HPLC. The isolated polypeptide of band A in Figure 3b was trypsinized and identified by microcapillary HPLC-tandem mass spectrometry. Peaks corresponding to *L. monocytogenes* flagellin peptides are indicated in Figure 4a. Five sequences were identified (Fig. 4a) that
25 correspond to flagellin, the product of the *flaA* gene of *L. monocytogenes* (Genbank Q02551). The location of these sequences within the protein is indicated in figure 4b.

Band B of Fig. 3b also is flagellin, which migrates as a doublet of approximately 30kDa on SDS-PAGE (Fig. 3b).

For analysis, bands A and B were excised from SDS-PAGE gels, dehydrated with acetonitrile, dried under reduced vacuum, and trypsin (12.5 ng/ μ L) was infused into the gel. The gel slice was allowed to incubate on ice for 45 min in the presence of trypsin and then excess trypsin removed and replaced with 50 mM ammonium bicarbonate and the gel slice incubated overnight at 37°C. Peptides were extracted by 3 washes with 5% acetic acid in 50% aqueous acetonitrile. The extractions were pooled and concentrated by vacuum centrifugation. The peptides were injected onto a C18 peptide trap cartridge (Michrom BioResources, Inc. Auburn, CA), desalted, and then injected onto a 75 μ m (internal diameter) x 10 cm micro-capillary HPLC column (Magic C18; 5- μ m packing; 100 A pore size; Michrom BioResources, Inc. Auburn, CA). The sample injection was made using a FAMOS autosampler (LCPackings, San Francisco, CA) coupled with an Agilent HP1100 Pump. Peptides were separated by a linear gradient of acetonitrile, and subjected to collision induced dissociation using an electrospray ionization-ion trap mass spectrometer (ESI-ITMS; ThermoQuest, San Jose, CA) in data-dependent mode with dynamic exclusion (Goodlett, et al. Anal. Chem. 72, 1112-1118 (2000)). Protein identification was accomplished by use of the SEQUEST computer program (Eng et al. J. Am. Soc. Mass. Spectrom. 5, 976-989 (1994)).

CHO cells expressing an NF- κ B luciferase reporter and TLR5 or TLR2 were stimulated with 100 μ l/ml

Listeria supernatant or 33 µg/ml purified *Salmonella* flagellin. Flagellin was purified from *Salmonella typhimurium* (TH4778 *fliB*- *fliC*+) by the procedure of Ibrahim et al., J. Clin. Microbiol. 22, 1040-1044 (1985).

5 As shown in Figure 4c, flagellin stimulated TLR5-expressing CHO cells, but not TLR2-expressing CHO cells. The mean and standard deviation of quadruplicate samples are indicated. CHO cells were transfected as described in above Examples with the addition of 0.1 µg of
10 pNeo/Tak, and stable populations of cells expressing the indicated TLR with the luciferase reporters were selected in 100 µg/ml G418. These cells were plated on 96-well plates at 100,000 cells/well, incubated overnight, and
processed in luciferase assays as described above.

15 The observation that flagellin is the TLR5 ligand also is supported by the finding that the flagellated bacteria, *L. monocytogenes* and *P. aeruginosa*, stimulate TLR5, while the TOP10 strain of *E. coli*, that
20 has lost its flagella, does not (Fig. 2b). Similarly, TLR5-stimulating activity was found in *B. subtilis*, *L. innocua*, *S. typhimurium* and *S. minnesota*, all flagellated bacteria, while non-flagellated bacteria such as *H. influenza*, did not activate TLR5.

25 Thus, the TLR5-stimulating activity purified from *L. monocytogenes* culture supernatants was identified as flagellin by tandem mass spectrometry.

EXAMPLE V

Flagellin Expression in Bacteria Reconstitutes
TLR5-Stimulating Activity

This Example shows that flagellin expression in
5 bacteria reconstitutes TLR-stimulating activity, and
deletion of flagellin genes abrogates TLR5-stimulating
activity.

To confirm that flagellin is the sole TLR5
ligand in bacteria, *E. coli* (TOP10) that secrete little
10 TLR5 activity (Fig. 2b) were transformed with the cDNA of
L. monocytogenes flagellin (*flaA*) under the control of an
inducible promoter. TLR-expressing CHO cells were
stimulated for 5 hours with *E. coli* culture supernatants
(67 µl/ml) in which expression of *L. monocytogenes*
15 flagellin was induced or repressed. In the
control sample, CHO cells were stimulated with
supernatants from induced *E. coli* containing the *L.*
monocytogenes flagellin gene cloned in the reverse
orientation. Supernatants of *E. coli* that were induced
20 to express *L. monocytogenes flaA* contained substantial
TLR5-stimulating activity (Fig. 5a), whereas supernatants
from *E. coli* in which expression was repressed, or from
E. coli expressing *flaA* in the reverse orientation,
contained little TLR5 activity in CHO cells expressing an
25 NF-κB luciferase reporter and TLR5 (Fig. 5a) or TLR2 (Fig
5b). CHO cells expressing an NF-κB luciferase reporter
and TLR5 (c) or TLR2 (d) were stimulated for 5 hours with
culture supernatants (100 µl/ml) from *S. typhimurium*
lacking one copy of flagellin (*FlhB- fliC+*) or both

copies of flagellin (*FliB*+ *FliC*+). Control is stimulation with LB medium. The mean and standard deviation of quadruplicate samples are indicated.

CHO cells were transfected with TLR2 and TLR5 expression plasmids as described above with the addition of 0.1 µg of pNeo/Tak, and stable populations of cells expressing the indicated TLR with the luciferase reporters were selected in 100 µg/ml G418. These cells were plated on 96-well plates at 100,000 cells/well, incubated overnight, and processed in luciferase assays as described above.

L. monocytogenes flagellin is not recognized by TLR2, since supernatants from *E. coli* expressing *flaA* did not show enhanced TLR2-dependent stimulation of CHO cells relative to supernatants from *E. coli* with repressed *flaA* expression (Fig. 5b). In addition to the experiments that demonstrate reconstitution of TLR5-stimulating activity by the expression of flagellin, a bacterium from which flagellin had been deleted was tested. It was observed that TLR5-stimulating activity was abrogated in the flagellin deleted strain. *S. typhimurium* possess two genes for flagellin, *fliB* and *fliC* (Fujita, J., J. Gen Microbiol. 76, 127-34 (1973)). Culture supernatants of *fliB*- *fliC*+ *S. typhimurium* contained TLR5-stimulating activity, while culture supernatants from *S. typhimurium* lacking both flagellins (*fliB*- *fliC*-) expressed no TLR5-stimulating activity (Fig. 5c). The lack of both flagellin genes had no effect on TLR2-stimulating activity (Fig. 5d). The observed TLR2-stimulating activity found in *S.*

typhimurium supernatants most likely was due to bacterial lipoproteins (Underhill, et al. Nature 401, 811-5 (1999); Brightbill et al., Science 285, 732-6 (1999)). These results indicate that flagellin is the sole

- 5 TLR5-stimulating activity present in *S. typhimurium* culture supernatant.

Thus, TLR5-stimulating activity was elicited by introducing the flagellin gene into a non-flagellated bacterium, and abrogated by deleting the flagellin genes
10 from a flagellated bacterium.

EXAMPLE VI

Flagellin-Induced System IL-6 Production in Mice

This example shows that TLR signaling is required for the *in vivo* immune response to flagellin.

- 15 To determine if TLR signaling is required for the *in vivo* immune response to flagellin, wild type mice and mice lacking a component of the TLR5 signal transduction pathway, MyD88, were injected with flagellin and systemic IL-6 production was monitored. MyD88 is an
20 adaptor protein required for TLR5-mediated signal transduction (Aderem A. and Ulevitch, R.J., Nature 406:782-787, (2000); Brightbill, H.D. and Modlin. R.L., Immunology 101:1-10, (2000)).

- MyD88^{-/-} mice (129/SvJ x C57B1/6 background)
25 were backcrossed for three generations with C57B1/6 mice (Adachi, O. et al. Immunity, 9:143-150 (1998)). Mice from the F₃ generation (MyD88^{-/-}, n=5) and littermate controls (MyD88^{+/+}, n=5) were injected i.p. with 30 µg

purified flagellin in 0.5 cc of saline. Blood was sampled at 0, 1, 2, 4 and 8 hours after injection, and IL-6 levels were determined by ELISA (Duoset, R&D Systems, Minneapolis, MN).

5 Figure 6 shows that flagellin induced systemic IL-6 within 2 h in wild type mice. By contrast, mice deficient in MyD88 were completely unresponsive to flagellin.

 Therefore, flagellin stimulates TLR5-mediated
10 responses *in vivo*.

 Throughout this application various publications have been referenced. The disclosures of these publications in their entireties are hereby incorporated by reference in this application in order to
15 more fully describe the state of the art to which this invention pertains.

 Although the invention has been described with reference to the disclosed embodiments, those skilled in the art will readily appreciate that the specific
20 experiments detailed are only illustrative of the invention. It should be understood that various modifications can be made without departing from the spirit of the invention.

What is claimed is:

1. An immunomodulatory flagellin peptide comprising at least about 10 amino acids of substantially the amino acid sequence GAVQNRFNSAIT (SEQ ID NO:2), or a
5 modification thereof, and having toll-like receptor 5 (TLR5) binding.
2. The flagellin peptide of claim 1, further comprising TLR5 stimulating activity.
- 10 3. The flagellin peptide of claim 1, further comprising an ADCC targeting molecule.
4. The flagellin peptide of claim 1, wherein said flagellin peptide comprises a peptide of
15 *S. Typhimurium*1 flagellin.
5. A method of inducing an antigen-specific immune response in an individual comprising, administering to an individual an immunogenic amount of a
20 vaccine, having an antigen and an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof.
- 25 6. The method of claim 5, wherein said antigen is selected from the group consisting of polypeptides, polysaccharides, pathologically aberrant cells and bacteria.

7. A method of inducing a TLR5-mediated response, comprising administering to a TLR5-containing cell an effective amount of an immunomodulatory flagellin peptide having at least about 10 amino acids of
5 substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof.

8. The method of claim 7, wherein said TLR5-mediated response is TLR5-induced modulation of cytokine
10 amount or activity.

9. The method of claim 7, wherein said TLR5-mediated response is TLR5-induced increase in an amount of a cytokine selected from the group consisting of TNF α , IL-1 and IL-6.

15 10. The method of claim 7, wherein said TLR5-mediated response is TLR5-induced NF- κ B activity.

11. A method of inducing an immune response in an individual having a pathological condition, comprising administering to said individual an immunogenic amount of
20 an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof

12. The method of claim 11, wherein said pathological condition is selected from the group
25 consisting of proliferative disease, autoimmune disease, infectious disease and inflammatory disease.

13. The method of claim 11, wherein said immunomodulatory flagellin peptide further comprises a heterologous amino acid sequence.

14. The method of claim 11, wherein said
5 immunomodulatory flagellin peptide further comprises an ADCC targeting molecule.

15. A method of modulating an immune response in an individual having a pathological condition,
10 comprising administering to said individual a combination of an immunogenic amount of an immunomodulatory flagellin peptide having at least about 10 amino acids of substantially the amino acid sequence of SEQ ID NO:2, or a modification thereof, and an immunomodulatory molecule.

15 16. The method of claim 15, wherein said immunomodulatory molecule is an antibody, cytokine or growth factor.

17. The method of claim 15, wherein said
20 immunomodulatory flagellin peptide further comprises a heterologous amino acid sequence.

18. The method of claim 15, wherein said immunomodulatory flagellin peptide further comprises an ADCC targeting molecule.

25 19. The method of claim 15, wherein said pathological condition is selected from the group consisting of proliferative disease, autoimmune disease, infectious disease and inflammatory disease.

20. A method of inducing an immune response in a human individual having a pathological condition, comprising administering to said individual an immunogenic amount of a flagellin polypeptide, or
5 modification thereof, wherein said flagellin polypeptide induces an immune response.

21. The flagellin polypeptide of claim 20, wherein said flagellin polypeptide comprises an
10 *S. Typhimurium*1 flagellin polypeptide.

22. The method of claim 20, wherein said pathological condition is selected from the group consisting of proliferative disease, autoimmune disease, infectious disease and inflammatory disease.

15 23. A screening composition comprising,
(a) a flagellin peptide of claim 1; and
(b) a TLR5 polypeptide or modification thereof, having a TLR5 activity.

20 24. The composition of claim 23, further comprising a detectably labeled flagellin peptide.

25 25. The composition of claim 23, wherein said TLR5 polypeptide comprises an amino acid sequence selected from the group consisting of SEQ ID NO:4 and
25 SEQ ID NO:6, or a modification or fragment thereof having a TLR5 activity.

26. A method of screening for a TLR5 ligand, agonist or antagonist, comprising:

- (a) contacting a TLR5 with a candidate compound in the presence of a flagellin polypeptide or
5 immunomodulatory flagellin peptide under conditions wherein binding of said flagellin polypeptide or immunomodulatory flagellin peptide to said TLR5 produces a predetermined signal;
- (b) determining the production of said
10 predetermined signal in the presence of said candidate compound; and
- (c) comparing said predetermined signal in the presence of said candidate compound with a predetermined signal in the absence of said candidate
15 compound, wherein a difference between said predetermined signals in the presence and absence of said candidate compound indicates that said compound is a TLR5 ligand, agonist or antagonist.

27. The method of claim 26, wherein said
20 predetermined signal is selected from the group consisting of polypeptide amount, polypeptide activity and transcriptional activity.

28. The method of claim 26, wherein said
predetermined signal is amount of a cytokine selected
25 from the group consisting of $\text{TNF}\alpha$, IL-1 and IL-6.

29. The method of claim 27, wherein said transcriptional activity is NF- κ B activity.

30. The method of claim 27, wherein said immunomodulatory flagellin peptide is a flagellin peptide of claim 1, or a modification thereof.

31. A bioassay for detecting bacterial
5 contamination in a sample comprising,
(a) contacting said sample with a TLR5 under conditions wherein binding of a flagellin polypeptide or fragment thereof in said sample to said TLR5 produces a predetermined signal;
10 (b) determining the production of said predetermined signal in the presence and absence of said sample; and
(c) comparing said predetermined signal in the presence of said sample with a predetermined
15 signal in the absence of said sample, wherein a difference between said predetermined signals in the presence and absence of said sample indicates that said sample contains flagellin.

32. The method of claim 31, wherein said
20 sample is a product for animal consumption.

33. The method of claim 31, wherein said predetermined signal is selected from the group consisting of polypeptide amount, polypeptide activity and transcriptional activity.

25 34. The method of claim 31, wherein said predetermined signal is NF- κ B activity.

35. The method of claim 31, wherein said predetermined signal is an amount a cytokine selected from the group consisting of TNF α , IL-1 and IL-6.

1/15

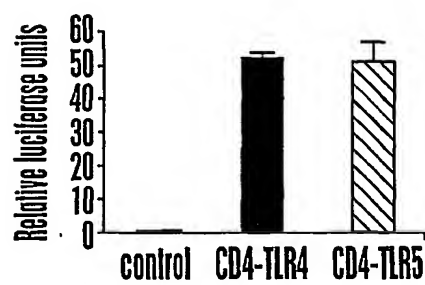


Figure 1A

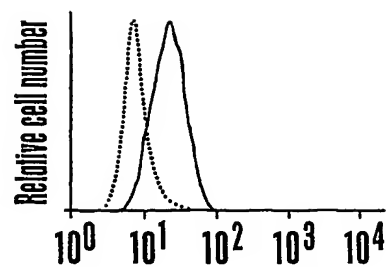


Figure 1B

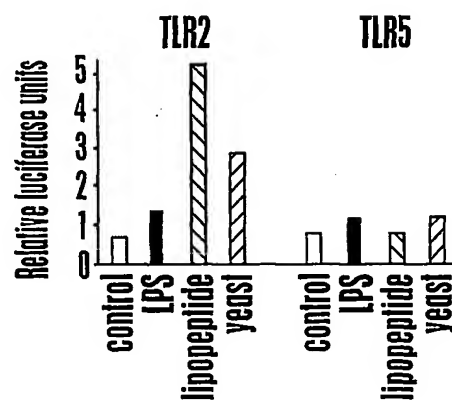


Figure 2A

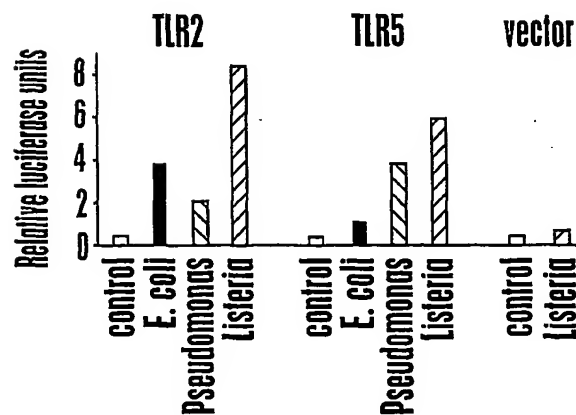


Figure 2B

Figure 3A

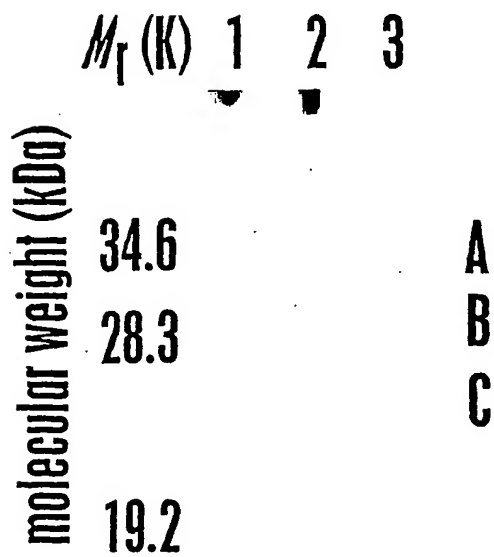
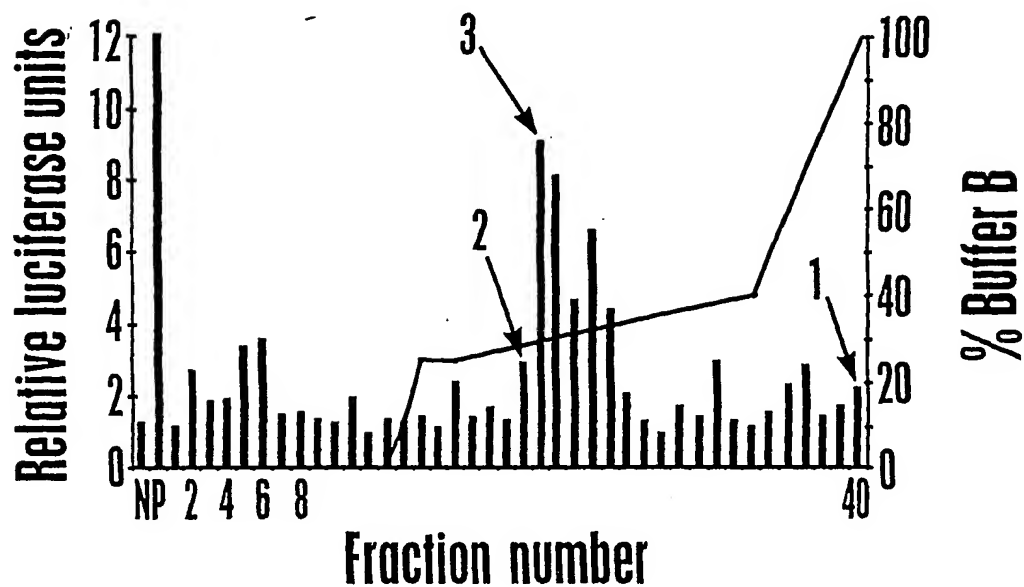


Figure 3B

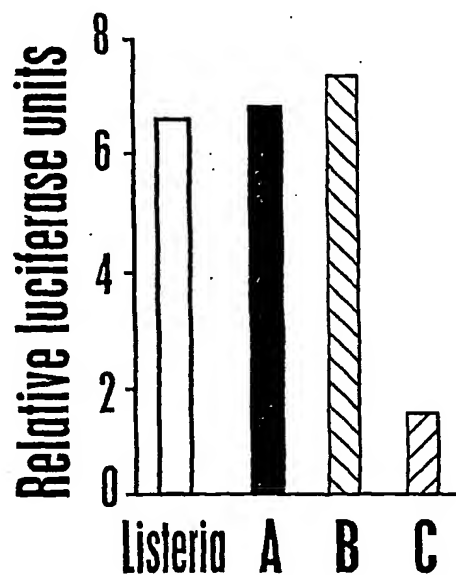


Figure 3C

3/15

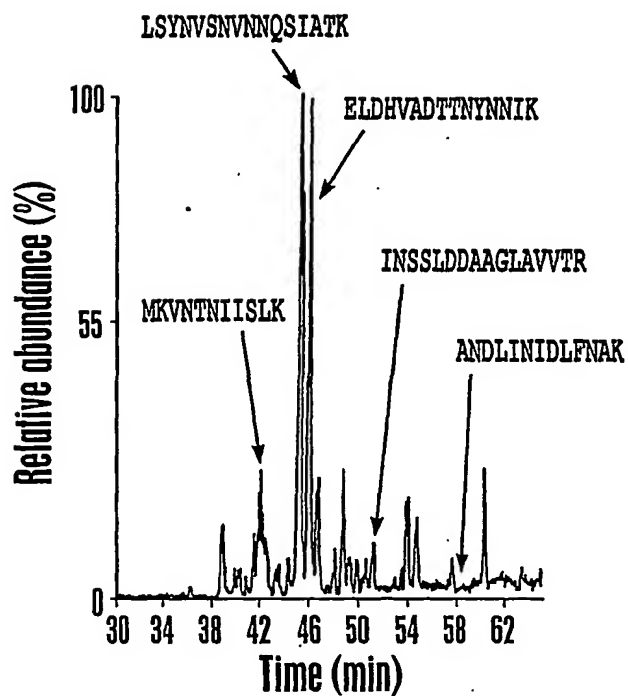
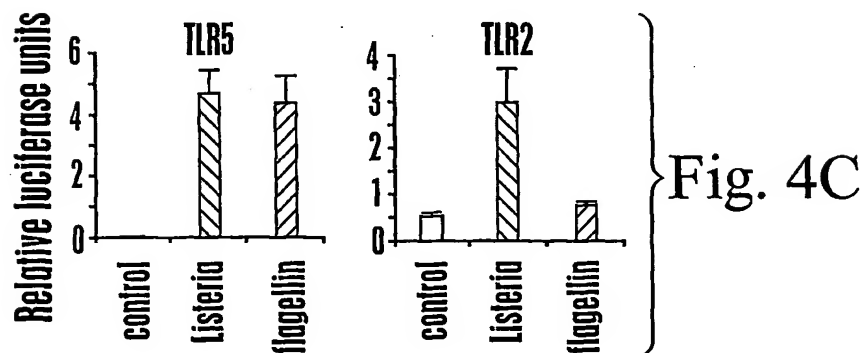


Fig. 4A

MKVNTNIISLK**TQ**EYLRKNEG**MTQ**AQERLASGKRINSSLD**D**
 AAGLAVVTRMNVKSTGLDAASKNSSMGIDLLQ**TAD**SALSSMS
 SILQRM**RQ**LAVQSSNGSFSDEDRKQY**TA**EFGSLIKELDHV**AD**
 TTN**NNIK**LLDQ**TAT**GAATQVSIQASDKANDLINIDLFNA**KG**
 LSAGTITLGSGSTVAGYSALS**VAD**ADSSQ**EATE**AIDELIN**NI**
 SNGRALLGAGMSRLSYNVSNNQSIATKASASSIEDADMA**A**
 EMSEMTKYKIL**TQ**TSISMLSQANQ**TPQ**ML**TQ**LIN**S**

Fig. 4B



4/15

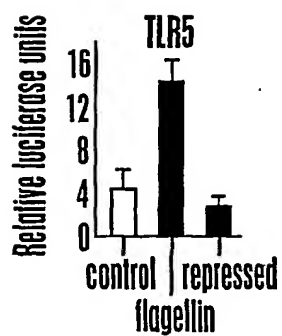


Fig. 5A

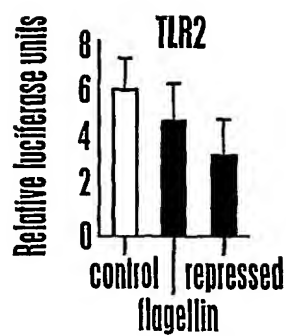


Fig. 5B

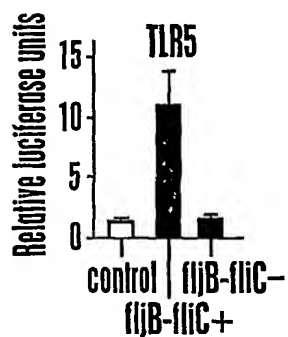


Fig. 5C

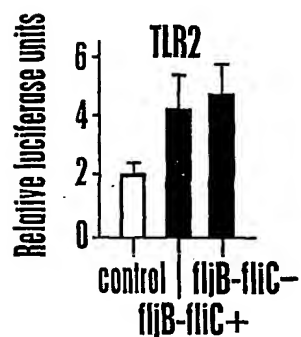


Fig. 5D

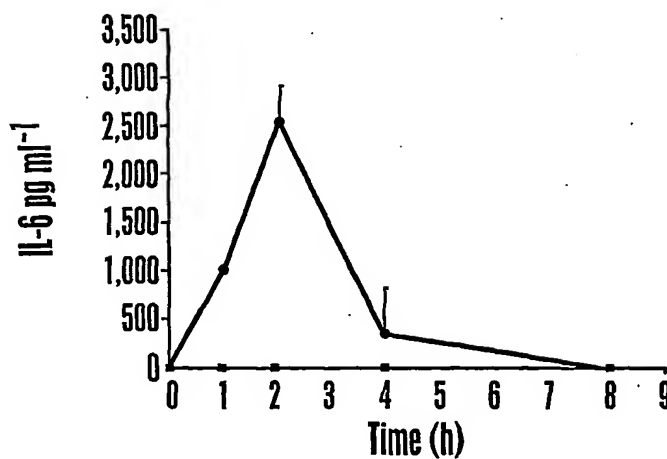


Fig. 6

5/15

C. JEJUNI	1	MGFRNTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOANT
H. PYLORI	1	MAFOVNTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
V. CHOLERA	1	MTINVTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
P. AERUGINOSA	1	MALVTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
R. SPHAEROIDES	1	-MTINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
P. MIRABILIS1	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
P. MIRABILIS2	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
S. TYPHIMURIUM2	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
S. TYPHIMURIUM1	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
S. MARCESENS	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
E. COLI	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
S. FLEXNERII	1	MAQVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
T. PALLIDUMA	1	--MTINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
T. PALLIDUMB	1	--MTINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
L. PNEUMOPHILA	1	--MTINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
B. BURGDOFFERII	1	--MTINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
B. SUBTILUS	1	--MRINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
C. DIFFICILE	1	--MRINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
R. MELILOTI	1	-MTSILTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
A. TUMEFACIENS	1	-MASILTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
R. LUPINI	1	-MASILTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
L. MONOCYTOGENES	1	--MKVINTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
B. CLARRIDGEIAE	1	MGTSILTNVAALNKAADLNKSLDASISRLSSGLRINSAADDAAGVATADTIRSOASS
CONSENSUS	1	m intNv al ag nl k g l slerlssGlrinSa ddaagmaia ri sqvrg

Figure 7AA

6/15

C. JEJUNI	61	LGQALSNQNDHIGLQVADKAMDEOLKILDTIKTKATOQAQD--GOSLKTRIMEQADINR
H. PYLORI	61	LGQALANTDQMGHLOVADKAMDEOLKILDTIKTKATOQAQD--GOTTESKKAQSDIVR
V. CHOLERAE	61	LDVAMRNANDGISIAQTAEGAMNESTSLORARDLQOSANG--TNSASERQAMNEEVA
P. AERUGINOSA	61	LNVAATRNANDGISIAQTAEGALQOSTNILORARDLSIOSANG--SNSDSERTAINGEAKQ
R. SPHAEROIDES	60	LNQATRNAQDCKNLVDTEGALHVEVSSMLORIRELAVOSND--TNTAADRGSHAAEGKQ
P. MIRABILIS1	61	LTOASRNANDGISIAQTEGALNEINNNLORIRELIVQAKNG--TNSNSDITSTQNEAKN
P. MIRABILIS2	61	LTOASRNANDGISISQTEGALNEINNNLORIRELIVQAKNG--TNSNSDINSTQNEAKQ
S. TYPHIMURIUM2	61	LTOASRNANDGISIAQTEGALNEINNNLORIRELAVOSANS--TNSOSDLSIQAEITQ
S. TYPHIMURIUM1	61	LTOASRNANDGISIAQTEGALNEINNNLORIRELAVOSANS--TNSOSDLSIQAEITQ
S. MARCESENS	61	LTOASRNANDGISIAQTEGALNEVNDMLQNRRLIVQANG--SNSDLSKSIQDEITQ
E. COLI	61	LTOASRNANDGISIAQTEGALSEINNNLORIRELIVQATG--TNSDSDLSIQDEEKS
S. FLEXNERII	61	LTOASRNANDGISIAQTEGALSEINNNLORIRELIVQASTG--TNSDSDLSIQDEEKS
T. PALLIDUMA	59	LNQASTNANSGINELQVTEAYLOETTDILORIRELAVQANG--IYSAEDRMQTOVEVSO
T. PALLIDUMB	59	LNQASTNANSGISIELQVATESYLOETTDILORIRELAVOSANG--IYSAEDRMQTOVEVSO
L. PNEUMOPHILA	59	LROAERNTEDEGMSIELQTAEGALQOSTNILORIRVLAVOTSNG--IYSNEDRQLQTOVEVSA
B. BURGDORFEREI	59	LSQASNTSKAINEIQTTEGALNEVEKMLVRMELAVOSNG--TYSADARGSTQDEITQ
B. SUBTILUS	59	LEMASKNSQDGISIELQTAEGALTETHAILORIRELIVQANGTQDKATDLOSIODEISA
C. DIFFICILE	59	LDOAGRNVDGISIELQTAEGALTEGNNILTRMRLAVOASNET--NSKDERAKIAGEEEO
R. MELILOTI	60	LSAVQDALGLCAKVDIASAGNESATIEWKEIKAKIQAATED-----GVDKAKIQDEITQ
A. TUMEFACIENS	60	LGAVSDALGMCAKVDIASAGNDAAIKWTDIKAKVVAKEQ-----GVDKTKIQDEVSO
R. LUPINI	60	LSAVQDAIGLCAKVDIASAGNDAAIKWTDIKAKVVAKEQ-----SADKTKIQDEVSO
L. MONOCYTOGENES	59	LDAASKNSSMCTDILQVADSALSSMSLORMROLAVOSSNG--SFSDEDRKQYTAFFGS
B. CLARRIDGEIAE	61	MSAIVDAINLEKEOVGIADDAIGLTKEADDHOKSMVSAREK-----GSDDIAKIQDSIG
CONSENSUS	61	l qatrnandgisilqtaegal e ilqiridl vqa ng tqe dr iq ei q

Figure 7AB

7/15

C. JEJUNI	119	IMEEDNTIANTTSFNGKQILSCNFINQEFQHCASSN-QTIKATIGATQSSKIGITRFETG
H. PYLORI	119	LLOGEDNIGNTTBYNGQALISQFTNKEFQHCAYSN-QSIKASIGSTTSKIGQVRIATG
V. CHOLERAE	119	LODEINRIIETTSTFGCKRLANGSFGEASFQHCSSSG-EAIIIGITSVRADDFR-----
P. AERUGINOSA	119	LOKEIDRIISNTTTEGCKRLIDSGFQVASFQHCSAAN-ETISVGIIDENSAESINGTYFKAD
R. SPHAEROIDES	118	LIAEINRVAESTTENGKVLIDCSFTGKQLQHCADSG-QTAINVDSAAATDICA-----
P. MIRABILIS1	119	VIDEINRISEOTQENGKVLISCEKSEMVICHTNDN-ETIKFNIDKVDNDTIGVASDKIF
P. MIRABILIS2	119	RIIDEINRVSEOTQENGKVLISCEKSEMTIQHTNDN-ETIEFNIDKVDNDTIGVASDKIF
S. TYPHIMURIUM2	119	RINEIDRVSGOTQENGKVLIA-QDNTLTIQHCANDG-ETIDIDEKQINSQTLGIDSLNVQ
S. TYPHIMURIUM1	119	RINEIDRVNGOTQENGKVLIA-QDNTLTIQHCANDG-ETIDIDEKQINSQTLGIDTLNVQ
S. MARCESENS	119	RIISEINRISEOTQENGKVLIS-SDQKLTIQHCANDG-ETTDIDEKKIDAKQIGMDTFDVT
E. COLI	119	RIIDEIDRVSGOTQENGKVLIS-KDGSMTIQHCANDG-ETITIDEKKIDSDTINAGFNIN
S. FLEXNERII	119	RIIDEIDRVSGOTQENGKVLIA-KDGSMTIQHCANDG-QTITIDEKKIDSDTIEGNGFNIN
T. PALLIDUMA	117	LIAEIDRIASSAQENGKVLITGRFSRTEG-----ENVIGGSMWFH
T. PALLIDUMB	117	LIAEIDRIASSAQENGKVLITGRFARETG-----ENTVTASMWFH
L. PNEUMOPHILA	117	LIIDEIDRIASSQAEINKFKLFEQFARGG-----R---VASMWFH
B. BURGDORFEREI	117	LTIDEINRIADQAOYNOMHMLSNKSASONVRTAEELGMQPAKINTPASLSGSOASWTLRVH
B. SUBTILUS	119	LTIDEIDGHSNRTEFNCKKILDSITYKVDATP-----ANQKNLVFQ
C. DIFFICILE	117	LRSEIDRIADSTKENGKVLISDDKKIALQVG-----AEAISNNVIEVS
R. MELILOTI	115	LKDQITSIAEAASESCENHLOADLSGGPVTKSVVGGFVRDSSGAVSVKKVDYSINTDTVL
A. TUMEFACIENS	115	LKDQIKSITGSASFCENHVLSSAN---ATKTVVSGFVRDAGGTVSUKTDTYALDANSML
R. LUPINI	115	LQEQKGIIVDSASFCENHVKCOLS-TTTTKSVVGSFVRE-GGTVSUKTIDYALNASKVL
L. MONOCYTOGENES	117	LKKEIDHIVADTININIKILDOTATGAATQVS-----IQASDKANDLINID
B. CLARRIDGEIAE	117	NKKNISNAVQSASFCENHLSNGGQTVGMAACYRREGTAIVYVDIIVGGSEINFGTIGSD
CONSENSUS	121	lmeeidria t fngmkil g qig v i v igl l

Figure 7BA

8/15

C. JEJUNI	178	GRISTSGEVQFTLKNYNGIDDFQFOKVVISTSVGTGLGALADEINKNADKTG----	VRAT
H. PYLORI	178	ALITASGDISLTFKQVDGVNDVTLESVKVSSSAGTGIGVLA EVT NKN SNRTG----	VKAY
V. CHOLERAE	171	MGGQSFLAEQPKTKEWGP-----	
P. AERUGINOSA	178	GGGAVTAATASGTVDIAIG-----	
R. SPHAEROIDES		-----	
P. MIRABILIS1	178	DTKTEKKGVTAA-----	
P. MIRABILIS2	178	DAKTEKKGVTAA-----	
S. TYPHIMURIUM2	177	KAYDVKDTAVTTKAYANNGTTLDVSGLDAAIKAATGGTNGTASVT-----	GGAVKFD
S. TYPHIMURIUM1	177	QKYKVSDTAATVTGYADTTIALDNS-----	TFKASATGLGGTDEKI-----DGD LKFD
S. MARCESENS	177	TKSAKAGAEIATG-----	
E. COLI	177	GEGETANTAATLKD MVGLKLDNTGVT TAGVNR YIADKAVASSTDILNAVAGVDGSKVSTE	
S. FLEXNERII	177	GGGAVANTAASKADLVAANATVVG NKYTVSAGYDAKASDLLAGVS---D---	GDTVQAT
T. PALLIDUMA	157	IGANMDQMR-----	VY-----
T. PALLIDUMB	157	IGANMDQTR-----	AY-----
L. PNEUMOPHILA	153	MGP NQNRER-----	FY-----
B. BURGDORFEREI	177	VGANQDEAIA-----	VN-----
B. SUBTILUS	159	IGANATQQIS-----	VN-----
C. DIFFICILE	160	LINTKGVLTT-----	RN-----
R. MELILOTI	175	FDTTGN---TGILDKVYN-----	
A. TUMEFACIENS	172	YTEG-----	
R. LUPINI	173	VDTRATG TKIGILD TAYTG-----	
L. MONOCYTOGENES	163	LFNAKGLSAG-----	
B. CLARRIDGEIAE	177	GTTDMSQGV LGGIFGTSKG-----	
CONSENSUS	181		

Figure 7BB

9/15

C. JEJUNI	234	FTVETRGLAAVRAGATSDTFAINGVKIGKVDYKGDANGALVAAINSVKDTTGVEASIDA
H. PYLORI	234	ASVITTSDEVAVQSGSLGNLTNGIHLGNIADIKKNSDGRLVAAINAVTSETGVEAYTDQ
V. CHOLERA	190	-----PTARDLKFEFTKK
P. AERUGINOSA	197	-----TTGGSANVVKVDM
R. SPHAEROIDES		-----
P. MIRABILIS1	191	-----AG-----VTDKAKINA
P. MIRABILIS2	191	-----DA-----IDANALGIS
S. TYPHIMURIUM2	230	ADNNKYFVTIGGFTGADAAKNG--DYEVNVATDGTVTLAAGATKTTMPAGATTKEVQEL
S. TYPHIMURIUM1	225	DTGKYYAKVTVTG--GTCKDG--YYEVSVDKTNGEVTAAVTPATVTTATALSCKMYSA
S. MARCESENS	190	-----T-----KITVDS--DA
E. COLI	237	ADVGFGAAPGTFPVEYTYHKDTNTYTASASVDATQLAAFLNPEAGGTTAATVSIGNGTTA
S. FLEXNERII	231	INNGFGTAASATNYKYDSASKS-YSFDTTASAADVQKYLTPGVGDTAKGTTTIDG---S
T. PALLIDUMA	169	-----IGTMTAVA
T. PALLIDUMB	169	-----IGTMTAAA
L. PNEUMOPHILA	165	-----IGTMTSKA
B. BURGDORFEREI	189	-----IYAANVAN
B. SUBTILUS	171	-----IEDMGADA
C. DIFFICILE	172	-----VNSANIDA
R. MELILOTI	190	-----VSQASVTLPVNV
A. TUMEFACIENS	176	-----T--
R. LUPINI	192	-----LNANTVTVDINK
L. MONOCYTOGENES	173	-----
B. CLARRIDGEIAE	196	-----DEGEDVVGKGIGA
CONSENSUS	241	

Figure 7CA

10/15

C. JEJUNI	294	NGQLLLTSREGRGIKIDGNIGGGAFINADMKENYGRSLVVRNDGKDILISGSNLSSAGFG
H. PYLORI	294	KGRINLRSIDGRGIEIK-----TDSVSNGPSALTMVNGGQDLTKGSTNYGRSLT
V. CHOLERA	203	DG----EAVVLDIIAKGD-----DIEELA-----TYINGQTD
P. AERUGINOSA	210	KGNETAEQAAKIAAAVND-----ANVCIG-----AFSDGDTI
R. SPHAEROIDES		-----
P. MIRABILIS1	202	AATLDMVSLVKEFNLDG-----KPVTDK-----FIVTKGGKD
P. MIRABILIS2	202	GSKKYVTGISVKEYKVDG-----KVSSDK-----VVLNDGSDD
S. TYPHIMURIUM2	288	KDTPAVVSADAKNALIAGGV-----DATDANGAELVKMSYTDKNGKTIEGGYALKAGDK
S. TYPHIMURIUM1	281	NPDSIAKAALTAAGVTG-----TASVVKMSYTDNNGKTIDGGLAVKVGDD
S. MARCESENS	199	T----KQADADVTGLAKG-----QTLVSG-----TDADGKSA
E. COLI	297	QEQKVIIAKDGSLLTAADDG-----AALYLDGTGNLSKTN-AGTDTQAKLS
S. FLEXNERII	287	-AQDVQISSDGKITASNG-----DKLYIDTTGRLTKNGSGASLTEASLS
T. PALLIDUMA	177	LG-----
T. PALLIDUMB	177	LG-----
L. PNEUMOPHILA	173	LK-----
B. BURGDORFEREI	197	LFSGEGAQAAQTAPVQEGA-----
B. SUBTILUS	179	LGIKEADG-----
C. DIFFICILE	180	MS-----
R. MELILOTI	202	NGTTSEYTVGAYNVDDLID-----ASATFDGDYANVGAGALAGDYVKVQG
A. TUMEFACIENS	177	-----
R. LUPINI	204	GGVITQASVRAYSTDEMLS-----LGAKVDGANSNVAVGGGSFVKVDGS
L. MONOCYTOGENES	173	-----
B. CLARRIDGEIAE	209	FSAAHATYKLEDTLRN-----AEADLAKAIKYGESPEDEPGKAI
CONSENSUS	301	

Figure 7CB

11/15

C. JEJUNI	354	ATQFISQASVSLRESKGQIDANIADAMGFGSANKGVVLGGYSSVSAY	MSAGSGFSSGSG
H. PYLORI	344	RLDAKSINNVVSAS-----DS-----Q-----	HGGFTAIGFGESQV
V. CHOLERA	232	LFKASVDQEGKIQ-----	TFVAEPNIEGNFN
P. AERUGINOSA	243	SYVSKAGKDGSGA-----	TSAVSGVVIADT
R. SPHAEROIDES		-----	
P. MIRABILIS1	235	YVATKSDFELDAT-----	GTK--LGLKASAT
P. MIRABILIS2	235	YIVSKSDFTLKSG-----	TTTGEVFTGSKT
S. TYPHIMURIUM2	342	YYAADYDEATGAI-----	KAKTTSYTAADGT
S. TYPHIMURIUM1	327	YYSATQDKDG-SI-----	SIDTTKYTADNGT
S. MARCESENS	227	YFIATKDDATGDV-----	AYTKAKVADDGKV
E. COLI	341	DLMANNANAKTVI-----	TT-DKGTFTANTT
S. FLEXNERII	330	TLAANNTKATTID-----	GGTSISFTGNST
T. PALLIDUMA	179	-----	VRNGVDESIMSIE
T. PALLIDUMB	179	-----	VRDVGDESILNID
L. PNEUMOPHILA	175	-----	IVKADGR-PIAIS
B. BURGENDORFEREI	216	QQEGAQQPAPVTA-----	PSQGGVNSPVNVT
B. SUBTILUS	187	---SIAALHSVND-----	IDVTKFADNAADT
C. DIFFICILE	182	-----	VS---GSI
R. MELILOTI	247	SWVKAVDVAATQGEVVYDD-----	GTTKWGVDTTVTGAPATNVA
A. TUMEFACIENS	177	--PGTIDANS-----	GILNATGATTTVG
R. LUPINI	249	WVKGSDAAAASITASTPVACK-----	FAAAYTAAEAGTAAAAGDAIVDRTNAGAGAV
L. MONOCYTOGENES	173	-----	TTTLGSGSTVAGYS
B. CLARRIDGEIAE	250	IEKAKQAVETAKTG-----	IKDGQEAATKAKG
CONSENSUS	361		II

Figure 7DA

12/15

C. JEJUNI	414	YSVSGSKNYSTGFANAIAISAASQLSTVYNVSAGSGFSSGSTLSQFA-----T-----
H. PYLORI	374	AETTVNLRDVTGNFNANVKSASGANYNAVIASGNQSLGSG-----
V. CHOLERA	258	ISGGLATELGLN-----
P. AERUGINOSA	269	GSTGVGTAAGVAPSA-----
R. SPHAEROIDES		-----
P. MIRABILIS1	259	TEFKVDAGKDVKTIN-----
P. MIRABILIS2	261	TKFTADAGKDVKVLN-----
S. TYPHIMURIUM2	368	TKTAANQLGGVDGKTEVVTIDGKTYNAS-----
S. TYPHIMURIUM1	352	SKTALNKLGGADGKTEVVTIDGKTYNAS-----
S. MARCESENS	253	TDSCTDAG-----
E. COLI	366	KFDGVDISVDASTFANAVKNETYTATVG--VTLPATYTVNNGTAASAYLVDGKVS KTP--
S. FLEXNERII	356	TPDTITYSVTGAKVDQAAFDKAVSTSGNNVDFTTAGYSVNGTTGAVTKGVD SVYVDNNEA
T. PALLIDUMA	192	--TADSAN-----
T. PALLIDUMB	192	--DPEKAN-----
L. PNEUMOPHILA	187	--SPGEAN-----
B. BURGDOFFERREI	242	--TTVDAN-----
B. SUBTILUS	210	--ADIGFD-----
C. DIFFICILE	187	--GTEAAS-----
R. MELILOTI	286	APASIAITIDITIAAQ-----
A. TUMEFACIENS	198	AKTYTQISVLD MNVG-----
R. LUPINI	302	NLTQSVLTMDVSSMS-----
L. MONOCYTOGENES	187	ALSVADAD-----
B. CLARRIDGEIAE	277	EFQTVLDGMTLADFTELKG-----
CONSENSUS	421	

Figure 7DB

13/15

C. JEJUNI	462	-----MKTTFAGVKDETAGVTTTLKGAMAVMDIAETATT
H. PYLORI	414	-----VTTLRGAMVVMDIAESAMK
V. CHOLERA	270	-----GGPGVKTTVQDIDITSVGGSONAVGIDDAIK
P. AERUGINOSA	284	-----TAFAKTNDTVAKIDISTAKLSRRACDRTTAIK
R. SPHAEROIDES		-----
P. MIRABILIS1	274	-----VKDDAATIDKAIN
P. MIRABILIS2	276	-----VKDDAATIDNAIS
S. TYPHIMURIUM2	396	-----KAAGHDFKAQPELAEAAKT TENPIQKIDAAIA
S. TYPHIMURIUM1	380	-----KAAGHDFKAEPELAEQAKT TENPIQKIDAAIA
S. MARCESENS	261	-----VKNPIATIDKAIIA
E. COLI	422	-----AEYFAQADGTITSGENAATSKAIYVSANGNLTTNTTSEATTNPIAAIDDAIA
S. FLEXNERII	416	LTTSDFVDFYLQDDGSVTNG----SGKAVYKDADGKLTDDAETKAATTADPKAIDDAIS
T. PALLIDUMA	198	-----KSICTIDDAIK
T. PALLIDUMB	198	-----RAICTIDDAIK
L. PNEUMOPHILA	193	-----DVIGLADAIET
B. BURGDORFEREI	248	-----TSLAKIENAIR
B. SUBTILUS	216	-----AOLKVVDEAIN
C. DIFFICILE	193	-----KMIVNIDSSIA
R. MELILOTI	301	-----AGNLDALTAGVDALT
A. TUMEFACIENS	213	-----TDDLDAIYSEITAIT
R. LUPINI	317	-----STDVGSYITGVEKAIT
L. MONOCYTOGENES	195	-----SSQATEAIDELIN
B. CLARRIDGEIAE	296	-----LGELHSDIQRMIMTSVONTVRDAIN
CONSENSUS	481	m id am

Figure 7EA

14/15

C. JEJUNI 495 NEDQIRADIGSVONOVSTINNIITVTOVNYKAESQIRVDFAESANYSKANITLAOSGS
 H. PYLORI 433 MIDKVRSDLGSTONOMISTINNIISITOVNYKAESQIRVDFAESANFNKNITLAOSGS
 V. CHOLERA 302 YDSDORADLGAKONRFSHSISNLSNIOENWASKSRKOTDFAKETTOITKSOILOOAGI
 P. AERUGINOSA 317 QIDASVPTSVAVONRFONTINNIKNIGENWSAARGRIEDTDFAEETANITKNOILOOAGI
 R. SPHAEROIDES -----
 P. MIRABILIS1 288 TIDESRSKLGAVONRFESTINNIINNTVNNLSASRSRIIDADYATEVSNMSRCOILLOOAGI
 P. MIRABILIS2 290 KIDESRSKLGAVONRFQSTINNIINNTVNNLSASRSRIIDADYATEVSNMSRCOILLOOAGI
 S. TYPHIMURIUM2 429 QVDALRSDLGAVONRFNSAITNIGNTVNNLSASRSRIEDSDYATEVSNMSRAOILLOOAGI
 S. TYPHIMURIUM1 413 QVDTLRSDLGAVONRFNSAITNIGNTVNNLSASRSRIEDSDYATEVSNMSRAOILLOOAGI
 S. MARCESENS 274 QVDGLRSSLGAVONRFDSVINNIINNTVNNLSASRSRIODADYATEVSNMSRANILLOOAGI
 E. COLI 476 SIDKFRSSLGAVONRFDSAVTNINNTTTNLSEASRIODADYATEVSNMSKAOILLOOAGI
 S. FLEXNERII 472 SIDKFRSSLGAVONRFDSAVTNINNTTTNLSEASRIODADYATEVSNMSKAOILLOOAGI
 T. PALLIDUMA 209 RINKORADLGAVONRFYTVGLDIAENIQAESRIIRDANIAKOMVEYTKOIVLOOAGI
 T. PALLIDUMB 209 KINKORADLGAVONRFYTVGLDIAENIQAESRIIRDANIAKOMVEYTKOIVLOOAGI
 L. PNEUMOPHILA 204 KINKORADLGAVONRFYTVGLDIAENIQAESRIIRDANIAKOMVEYTKOIVLOOAGI
 B. BURGDORFEREI 259 MISDORANLGAVONRFESIKDSTEYAIENIKASYAQIKDAMTDEIVAAITNSILLOOAGI
 B. SUBTILUS 227 QVSSORAKLGAVONRFLEHTINNIISASGENITAEESRIIRDANIAKOMVEYTKOIVLOOAGI
 C. DIFFICILE 204 DINSARALLGAVONRFLESTONNIINNTVNNLSASRSRIODADYATEVSNMSKAOILLOOAGI
 R. MELLITI 317 DMTSAAASLGSSISRIIDLOQDFNKLSDSIDSGVGRVADAMNEESTKIKALOTOOLAI
 A. TUMEFACIENS 229 KMTSAGAKLGSSISRIIDLOQSGFADKLSDEKGVGRVADAMNEESTKIKALOTOOLAI
 R. LUPINI 333 SETSAGAKLGSSISRIIDLOQDFNKLSDSIDSGVGRVADAMNEESTKIKALOTOOLAI
 L. MONOCYTOGENES 209 NISNGRALLGAVONRFISYNVSNVNNQSIATKASASSIEDADMAEMSEMIKYKITOTSI
 B. CLARRIDGEIAE 321 VTLTAGSKLGAVONRFINIOENFKKLLDNIEVIGAGVADAMNEESTKIKALOTOOLAI
 CONSENSUS 541 l ra lgavqnrvd i nl enl aa sri dad a evtnlsk qilqq gs

Figure 7EB

15/15

C. JEJUNI	555	YAAQANSVHONVLRILIQ--
H. PYLORI	493	YAMSOANTVQONILRLIT--
V. CHOLERAE	362	SLAQAKOLENSAISLIQ--
P. AERUGINOSA	377	AMLAQANQEPQSVLSLLR--
R. SPHAEROIDES		-----
P. MIRABILIS1	348	SLAQANQEPQVLSLLR--
P. MIRABILIS2	350	AMLAQANQEPQVLSLLR--
S. TYPHIMURIUM2	489	SLAQANQEPQVLSLLR--
S. TYPHIMURIUM1	473	SLAQANQEPQVLSLLR--
S. MARCESENS	334	SLAQANQSTQVLSLLR--
E. COLI	536	SLAKANQEPQVLSLQOG-
S. FLEXNERII	532	SLAKANQEPQVLSLQOG-
T. PALLIDUMA	269	AMLAQANTSAQSILSLR--
T. PALLIDUMB	269	AMLAQANCATQSVLSLLR--
L. PNEUMOPHILA	264	AMLRANMKENSVLRLIHI
B. BURGDOFFEREI	319	AMLAQANQEPQVLSLLR--
B. SUBTILUS	287	AMLAQANQEPQVLSLLR--
C. DIFFICILE	264	SLAQANQEPQVLSLLR--
R. MELIOTI	377	QALSTANSDSQVLSLLR--
A. TUMEFACIENS	289	QALSTANSDSQVLSLLR--
R. LUPINI	393	QALSTANSDSQVLSLLR--
L. MONOCYTOGENES	269	SLAQANQEPQVLSLLR--
B. CLARRIDGEIAE	381	QALSTANQGSQVLSLLR--
CONSENSUS	601	ilaqanq pgnvlsllr

Figure 7F

SEQUENCE LISTING

<110> THE INSTITUTE FOR SYSTEMS BIOLOGY

<120> TOLL-LIKE RECEPTOR 5 LIGANDS AND METHODS
OF USE

<130> FP-IS 4860

<150> US 60/285,477

<151> 2001-04-20

<160> 8

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 36

<212> DNA

<213> Salmonella typhimurium

<220>

<221> CDS

<222> (1)...(36)

<400> 1

ggt gcg gta cag aac cgt ttc aac tcc gct att acc

36

Gly Ala Val Gln Asn Arg Phe Asn Ser Ala Ile Thr

1

5

10

<210> 2

<211> 12

<212> PRT

<213> Salmonella typhimurium

<400> 2

Gly Ala Val Gln Asn Arg Phe Asn Ser Ala Ile Thr

1

5

10

<210> 3

<211> 4286

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (999)...(3575)

<400> 3

ttgaaatctc acagcccggt tggttgcagt gaccacttc gttgaacata ttcttcctaa 60

tcctagtact ttcaatttgc tctattccct ggtgtctatg catttaaadc gactatgggg 120
 ccattcttcc ttgaaccacc acagaagaca ttagctctct gggatccttg ttaatttttt 180
 ctcctcttac atagcaccta cgcttggaac atatgccaga cacatctgtg agacacccct 240
 tgccgctgca gctcatggat ggatgctgag ttccccacg caccacactt cagcaggtgg 300
 gtgtatttct gcttcacatt atactcccac acggccatgc atgtcaggca tggagcaggc 360
 tcataaccca cttaattaag gtgatcatat cagatccttt atcaagatgc atagagtgtc 420
 cagtgcctgt actatgatct cggatctttg ggagatgggc tagatagagt ctgggacaga 480
 atacagcaga gaaaccgata tgtttattgt ccgatcatca gctaagcttc tgggagctag 540
 gaatggggct ccttgatga acagaagtaa aaatgcctcg tctttatgac tttcaacttc 600
 cctcagcagg tctggaatgg gtgaacaaac actgcctgcg tgggtgataa atagcctctt 660
 tttgctgctt gtttgcctgt tttatggttc tgggagggaa cctagaacct agcacatgct 720
 agacaagtc tctagcactg agctatctcc ccagcttgga tgaaatatct gtaaagtact 780
 ggtgcccgctg tgtaaaatat gcaccattaa gtgttcaaga agaaaagact gggcatttct 840
 gttccaccaa gacaagaaga atctgccagc agaattgttg cgcagtcatt tgagcaaagg 900
 ggtccaaggg acagtaccct ccagtgtgg ggacccatgt gccgagcctc aggtgtgtgat 960
 gtgggtgtgt ttttaattct ctcttttccc ataggatc atg gca tgt caa ctt gac 1016

Met Ala.Cys Gln Leu Asp

1

5

ttg ctc ata ggt gtg atc ttc atg gcc agc ccc gtg ttg gta ata tct 1064
 Leu Leu Ile Gly Val Ile Phe Met Ala Ser Pro Val Leu Val Ile Ser
 10 15 20

ccc tgt tct tca gac ggc agg ata gcc ttt ttc cga ggc tgt aac ctc 1112
 Pro Cys Ser Ser Asp Gly Arg Ile Ala Phe Phe Arg Gly Cys Asn Leu
 25 30 35

acc cag att ccc tgg atc ctc aat act acc act gag agg ctc ctg ctc 1160
 Thr Gln Ile Pro Trp Ile Leu Asn Thr Thr Thr Glu Arg Leu Leu Leu
 40 45 50

agc ttc aac tat atc agt atg gtg gtt gcc aca tca ttt cca ctc ctg 1208
 Ser Phe Asn Tyr Ile Ser Met Val Val Ala Thr Ser Phe Pro Leu Leu
 55 60 65 70

gag cgg ctc cag ttg ctg gag ctg ggg acc cag tat gct aac ttg acc 1256
 Glu Arg Leu Gln Leu Leu Glu Leu Gly Thr Gln Tyr Ala Asn Leu Thr
 75 80 85

att ggt cca ggg gct ttc aga aac ctg ccc aat ctt agg atc ttg gac 1304
 Ile Gly Pro Gly Ala Phe Arg Asn Leu Pro Asn Leu Arg Ile Leu Asp
 90 95 100

ttg ggc caa agc cag atc gaa gtc ttg aat cga gat gcc ttt caa ggt 1352
 Leu Gly Gln Ser Gln Ile Glu Val Leu Asn Arg Asp Ala Phe Gln Gly
 105 110 115

ctg ccc cat ctc ttg gaa ctt cgg ctg ttt tcc tgt gga ctc tcc agt 1400
 Leu Pro His Leu Leu Glu Leu Arg Leu Phe Ser Cys Gly Leu Ser Ser
 120 125 130

gct gtg tta agt gac ggt tac ttc aga aat cta tat tca tta gct cgc 1448
 Ala Val Leu Ser Asp Gly Tyr Phe Arg Asn Leu Tyr Ser Leu Ala Arg
 135 140 145 150

WO 02/085933 PCT/US01/22978

tta gac cta tct ggc aac cag att cac agc ctc cgc ctc cat tct tca	1496
Leu Asp Leu Ser Gly Asn Gln Ile His Ser Leu Arg Leu His Ser Ser	
155 160 165	
ttc cgg gaa ctg aat tcc tta agc gac gta aat ttt gct ttc aac caa	1544
Phe Arg Glu Leu Asn Ser Leu Ser Asp Val Asn Phe Ala Phe Asn Gln	
170 175 180	
ata ttc act ata tgt gaa gat gaa ctc gag cct ctg cag ggc aaa aca	1592
Ile Phe Thr Ile Cys Glu Asp Glu Leu Glu Pro Leu Gln Gly Lys Thr	
185 190 195	
ctg tct ttc ttt ggc ctc aaa tta act aag ctg ttc agc aga gtc tct	1640
Leu Ser Phe Phe Gly Leu Lys Leu Thr Lys Leu Phe Ser Arg Val Ser	
200 205 210	
gtg ggc tgg gag aca tgc agg aac ccc ttc aga ggc gtg agg cta gaa	1688
Val Gly Trp Glu Thr Cys Arg Asn Pro Phe Arg Gly Val Arg Leu Glu	
215 220 225 230	
act cta gat ctt tct gaa aat ggc tgg acg gtg gac atc aca agg aac	1736
Thr Leu Asp Leu Ser Glu Asn Gly Trp Thr Val Asp Ile Thr Arg Asn	
235 240 245	
ttc agc aac atc atc cag gga agc cag att tcc tct ttg att ctt aaa	1784
Phe Ser Asn Ile Ile Gln Gly Ser Gln Ile Ser Ser Leu Ile Leu Lys	
250 255 260	
cac cac atc atg ggt cct ggc ttt ggc ttc cag aac atc aga gat cct	1832
His His Ile Met Gly Pro Gly Phe Gly Phe Gln Asn Ile Arg Asp Pro	
265 270 275	
gac cag agc aca ttt gcc agc ctg gcc aga agt tcg gtg ctg caa ctg	1880
Asp Gln Ser Thr Phe Ala Ser Leu Ala Arg Ser Ser Val Leu Gln Leu	
280 285 290	
gac ctt tcg cac ggc ttt atc ttc tcc ttg aat cct cga ctg ttt ggg	1928
Asp Leu Ser His Gly Phe Ile Phe Ser Leu Asn Pro Arg Leu Phe Gly	
295 300 305 310	
aca ctg aag gat ttg aag atg ctg aac ctt gcc ttc aac aag ata aac	1976
Thr Leu Lys Asp Leu Lys Met Leu Asn Leu Ala Phe Asn Lys Ile Asn	
315 320 325	
aag att gga gag aat gcc ttt tat ggg ctt gac agc ctc cag gtt ctc	2024
Lys Ile Gly Glu Asn Ala Phe Tyr Gly Leu Asp Ser Leu Gln Val Leu	
330 335 340	
aat cta tcc tat aat ctt ttg ggg gaa ctc tat aat tcc aac ttc tat	2072
Asn Leu Ser Tyr Asn Leu Leu Gly Glu Leu Tyr Asn Ser Asn Phe Tyr	
345 350 355	
ggg ctt cct aga gta gcc tac gtt gac ctt caa agg aac cac att ggg	2120
Gly Leu Pro Arg Val Ala Tyr Val Asp Leu Gln Arg Asn His Ile Gly	
360 365 370	

atc att caa gac caa aca ttc aga tta tta aaa acg tta caa acc tta	2168
Ile Ile Gln Asp Gln Thr Phe Arg Leu Leu Lys Thr Leu Gln Thr Leu	
375 380 385 390	
gat ctc cgt gac aat gct ctt aag gcc att ggt ttt att cca agc ata	2216
Asp Leu Arg Asp Asn Ala Leu Lys Ala Ile Gly Phe Ile Pro Ser Ile	
395 400 405	
cag atg gtc ctc ctg gga ggc aat aag ctg gtc cat ttg cca cac atc	2264
Gln Met Val Leu Leu Gly Gly Asn Lys Leu Val His Leu Pro His Ile	
410 415 420	
cac ttt act gcc aac ttc cta gag tta tct gaa aac agg cta gaa aac	2312
His Phe Thr Ala Asn Phe Leu Glu Leu Ser Glu Asn Arg Leu Glu Asn	
425 430 435	
ctg tcc gac ctc tac ttc ctc ctg cga gtc ccc cag ctc cag ttt ctc	2360
Leu Ser Asp Leu Tyr Phe Leu Leu Arg Val Pro Gln Leu Gln Phe Leu	
440 445 450	
atc ttg aat cag aat cgc ctt tcg tca tgc aag gca gcc cac act ccc	2408
Ile Leu Asn Gln Asn Arg Leu Ser Ser Cys Lys Ala Ala His Thr Pro	
455 460 465 470	
tcg gag aac cca agc tta gaa cag ctt ttc ctt aca gag aat atg ctg	2456
Ser Glu Asn Pro Ser Leu Glu Gln Leu Phe Leu Thr Glu Asn Met Leu	
475 480 485	
cag ctg gcc tgg gag acc ggc ctc tgt tgg gat gtt ttt caa ggc ctt	2504
Gln Leu Ala Trp Glu Thr Gly Leu Cys Trp Asp Val Phe Gln Gly Leu	
490 495 500	
tcc cgc ctc cag att ctt tac ctg agt aat aac tac ctt aat ttc ctt	2552
Ser Arg Leu Gln Ile Leu Tyr Leu Ser Asn Asn Tyr Leu Asn Phe Leu	
505 510 515	
cca cct ggg ata ttt aac gac ctg gtt gca tta cgg atg ctt agt ctt	2600
Pro Pro Gly Ile Phe Asn Asp Leu Val Ala Leu Arg Met Leu Ser Leu	
520 525 530	
agt gct aac aag ctg acc gtg ctc tct ccg ggc agt tta cct gct aat	2648
Ser Ala Asn Lys Leu Thr Val Leu Ser Pro Gly Ser Leu Pro Ala Asn	
535 540 545 550	
tta gag att ctc gac ata tct aga aat cag ctt ttg tgt cct gac cct	2696
Leu Glu Ile Leu Asp Ile Ser Arg Asn Gln Leu Leu Cys Pro Asp Pro	
555 560 565	
gct ttg ttt tct tcg ctt cgt gtt ttg gac ata act cat aac gag ttc	2744
Ala Leu Phe Ser Ser Leu Arg Val Leu Asp Ile Thr His Asn Glu Phe	
570 575 580	
gtc tgc aac tgt gaa ctt agc act ttt atc tcc tgg ctc aac caa acc	2792
Val Cys Asn Cys Glu Leu Ser Thr Phe Ile Ser Trp Leu Asn Gln Thr	

585	590	595	
aac gtc acc ctg ttc ggc tct cct gca gac gtg tat tgc atg tac cct			2840
Asn Val Thr Leu Phe Gly Ser Pro Ala Asp Val Tyr Cys Met Tyr Pro			
600	605	610	
aac tca ctg cta ggg ggc tcc ctc tac aac ata tcc acc gaa gac tgc			2888
Asn Ser Leu Leu Gly Gly Ser Leu Tyr Asn Ile Ser Thr Glu Asp Cys			
615	620	625	630
gat gaa gag gaa gcc atg cgg tcc cta aag ttt tcc ctt ttc atc ctg			2936
Asp Glu Glu Glu Ala Met Arg Ser Leu Lys Phe Ser Leu Phe Ile Leu			
635	640	645	
tgc acg gtc act ttg act cta ttc ctc gtc atc acc ctt gta gtc ata			2984
Cys Thr Val Thr Leu Thr Leu Phe Leu Val Ile Thr Leu Val Val Ile			
650	655	660	
aag ttc cgg gga atc tgt ttc ctg tgc tat aag acc atc cag aag ctg			3032
Lys Phe Arg Gly Ile Cys Phe Leu Cys Tyr Lys Thr Ile Gln Lys Leu			
665	670	675	
gtg ttc aag gac aag gtc tgg agt ttg gaa cct ggt gca tat aga tat			3080
Val Phe Lys Asp Lys Val Trp Ser Leu Glu Pro Gly Ala Tyr Arg Tyr			
680	685	690	
gat gcc tac ttc tgc ttc agc agc aaa gac ttt gaa tgg gca cag aat			3128
Asp Ala Tyr Phe Cys Phe Ser Ser Lys Asp Phe Glu Trp Ala Gln Asn			
695	700	705	710
gct ttg ctc aaa cac ctg gat gct cac tac agt tcc cga aac agg ctc			3176
Ala Leu Leu Lys His Leu Asp Ala His Tyr Ser Ser Arg Asn Arg Leu			
715	720	725	
agg cta tgc ttt gaa gaa aga gac ttc att ccg ggg gaa aac cat atc			3224
Arg Leu Cys Phe Glu Glu Arg Asp Phe Ile Pro Gly Glu Asn His Ile			
730	735	740	
tcc aac atc cag gcg gct gtc tgg ggc agc agg aag acg gtg tgt cta			3272
Ser Asn Ile Gln Ala Ala Val Trp Gly Ser Arg Lys Thr Val Cys Leu			
745	750	755	
gtg agc aga cac ttc ctg aag gat ggt tgg tgc ctg gag gcc ttc agg			3320
Val Ser Arg His Phe Leu Lys Asp Gly Trp Cys Leu Glu Ala Phe Arg			
760	765	770	
tat gcc cag agc cgg agt ctg tct gac ctc aag agc att ctc atc gtg			3368
Tyr Ala Gln Ser Arg Ser Leu Ser Asp Leu Lys Ser Ile Leu Ile Val			
775	780	785	790
gtg gtg gtg gga tcg ctg tcc cag tat cag ctg atg aga cat gag acc			3416
Val Val Val Gly Ser Leu Ser Gln Tyr Gln Leu Met Arg His Glu Thr			
795	800	805	
atc aga ggg ttt ctg caa aag caa cag tac ttg agg tgg cct gaa gac			3464

Ile Arg Gly Phe Leu Gln Lys Gln Gln Tyr Leu Arg Trp Pro Glu Asp
 810 815 820

ctc cag gat gtt ggc tgg ttt ctc gat aaa ctc tcc gga tgc att cta 3512
 Leu Gln Asp Val Gly Trp Phe Leu Asp Lys Leu Ser Gly Cys Ile Leu
 825 830 835

aag gaa gaa aaa gga aag aaa aga agc agt tcc atc cag ttg cga acc 3560
 Lys Glu Glu Lys Gly Lys Lys Arg Ser Ser Ser Ile Gln Leu Arg Thr
 840 845 850

ata gca acc att tcc tagcaggagc gcctcctagc agaagtgcaa gcatcgtaga 3615
 Ile Ala Thr Ile Ser
 855

taactctcca cgctttatcc gcacagccgc tgggggtcct tccctggagt catttttctg 3675
 acaatgaaaa caacaccaat ctcttgattt ttcattgtcaa cagggagctt tgtcttcact 3735
 gttttccaaa tggaagtaa gaggtccaga aagctgcctc taagggctct cacctgccat 3795
 tgatgtcctt tcaggcccaa tgacatggtt tccctccatc ctattgcgta ctgtctgcta 3855
 cccaggtggc aagagcacct tgggagaagt tacaggcagc ttcattgctt ctgtgctgtt 3915
 cagttcaaaa gcaggtgcct tgagaatcct gaattcaagc actctgtaga acatggacag 3975
 acaagatggg tccttctctg gccataggca tgagggccag ttgctgagga ctgctctcac 4035
 tacacctaag tgcacaagtg ataagaagt ggacagatag acagatagca gcagtcccat 4095
 tgctgtagcc agaatgcact tatttcctgt tctgaccctg cagggccagc ttttggggac 4155
 cacagccatg ttctgcacgg gacctctcaa cctggcattc atgcccttcc acgacttagc 4215
 accggcctgc ccttctttct tccccacaac tatacaagag ctgttgcaac cactgaaaaa 4275
 aaaaaaaaa a 4286

<210> 4

<211> 859

<212> PRT

<213> Mus musculus

<400> 4

Met Ala Cys Gln Leu Asp Leu Leu Ile Gly Val Ile Phe Met Ala Ser
 1 5 10 15

Pro Val Leu Val Ile Ser Pro Cys Ser Ser Asp Gly Arg Ile Ala Phe
 20 25 30

Phe Arg Gly Cys Asn Leu Thr Gln Ile Pro Trp Ile Leu Asn Thr Thr
 35 40 45

Thr Glu Arg Leu Leu Leu Ser Phe Asn Tyr Ile Ser Met Val Val Ala
 50 55 60

Thr Ser Phe Pro Leu Leu Glu Arg Leu Gln Leu Leu Glu Leu Gly Thr
 65 70 75 80

Gln Tyr Ala Asn Leu Thr Ile Gly Pro Gly Ala Phe Arg Asn Leu Pro
 85 90 95

Asn Leu Arg Ile Leu Asp Leu Gly Gln Ser Gln Ile Glu Val Leu Asn
 100 105 110

Arg Asp Ala Phe Gln Gly Leu Pro His Leu Leu Glu Leu Arg Leu Phe
 115 120 125

Ser Cys Gly Leu Ser Ser Ala Val Leu Ser Asp Gly Tyr Phe Arg Asn
 130 135 140

Leu Tyr Ser Leu Ala Arg Leu Asp Leu Ser Gly Asn Gln Ile His Ser
 145 150 155 160

Leu Arg Leu His Ser Ser Phe Arg Glu Leu Asn Ser Leu Ser Asp Val

- 7 -

Val Tyr Cys Met Tyr Pro Asn Ser Leu Leu Gly Gly Ser Leu Tyr Asn
 610 615 620
 Ile Ser Thr Glu Asp Cys Asp Glu Glu Glu Ala Met Arg Ser Leu Lys
 625 630 635 640
 Phe Ser Leu Phe Ile Leu Cys Thr Val Thr Leu Thr Leu Phe Leu Val
 645 650 655
 Ile Thr Leu Val Val Ile Lys Phe Arg Gly Ile Cys Phe Leu Cys Tyr
 660 665 670
 Lys Thr Ile Gln Lys Leu Val Phe Lys Asp Lys Val Trp Ser Leu Glu
 675 680 685
 Pro Gly Ala Tyr Arg Tyr Asp Ala Tyr Phe Cys Phe Ser Ser Lys Asp
 690 695 700
 Phe Glu Trp Ala Gln Asn Ala Leu Leu Lys His Leu Asp Ala His Tyr
 705 710 715 720
 Ser Ser Arg Asn Arg Leu Arg Leu Cys Phe Glu Glu Arg Asp Phe Ile
 725 730 735
 Pro Gly Glu Asn His Ile Ser Asn Ile Gln Ala Ala Val Trp Gly Ser
 740 745 750
 Arg Lys Thr Val Cys Leu Val Ser Arg His Phe Leu Lys Asp Gly Trp
 755 760 765
 Cys Leu Glu Ala Phe Arg Tyr Ala Gln Ser Arg Ser Leu Ser Asp Leu
 770 775 780
 Lys Ser Ile Leu Ile Val Val Val Val Gly Ser Leu Ser Gln Tyr Gln
 785 790 795 800
 Leu Met Arg His Glu Thr Ile Arg Gly Phe Leu Gln Lys Gln Gln Tyr
 805 810 815
 Leu Arg Trp Pro Glu Asp Leu Gln Asp Val Gly Trp Phe Leu Asp Lys
 820 825 830
 Leu Ser Gly Cys Ile Leu Lys Glu Lys Gly Lys Lys Arg Ser Ser
 835 840 845
 Ser Ile Gln Leu Arg Thr Ile Ala Thr Ile Ser
 850 855

<210> 5

<211> 3431

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (704)...(3277)

<400> 5

ggcttatagg gctcgagcgg ccgcccgggc aggtatagaa ttcagcggcc gctgaattct 60
 agggttttca ggagcccagag cgagggcgcc gcttttgcgt ccgggaggag ccaaccgtgg 120
 cgcaggcggc gcggggaggc gtcccagagt ctcaactctgc cgcccaggct ggactgcagt 180
 gacacaatct cggctgactg caaccactgc ctccagggtt caagcgattc tcttgccctca 240
 gcctcccaag tagctgggat tacagattga tgttcattgtt cctggcacta ctacaagatt 300
 catactcctg atgctactga caacgtggct tctccacagt caccaaacca gggatgctat 360
 actggacttc cctactctca tctgctccag cccctgacc ttatagttgc ccagctttcc 420
 tggcaattga ctttgcccat caatacacag gatttagcat ccaggaaga tgcggagcc 480
 tcagatgtta attttctaata tgagaatgtt ggcgctgtcc gaacctggag acagaaaaac 540
 aaaaagtcct ttctcctgat tcacaaaaaa ataaaatact gactaccatc actgtgatga 600
 gattcctata gtctcaggaa ctgaagtctt taaacaacca gggaccctct gccctagaa 660

taagaacata ctagaagtcc cttctgctag gacaacgagg atc atg gga gac cac	715
Met Gly Asp His	
1	
ctg gac ctt ctc cta gga gtg gtg ctc atg gcc ggt cct gtg ttt gga	763
Leu Asp Leu Leu Leu Gly Val Val Leu Met Ala Gly Pro Val Phe Gly	
5 10 15 20	
att cct tcc tgc tcc ttt gat ggc cga ata gcc ttt tat cgt ttc tgc	811
Ile Pro Ser Cys Ser Phe Asp Gly Arg Ile Ala Phe Tyr Arg Phe Cys	
25 30 35	
aac ctc acc cag gtc ccc cag gtc ctc aac acc act gag agg ctc ctg	859
Asn Leu Thr Gln Val Pro Gln Val Leu Asn Thr Thr Glu Arg Leu Leu	
40 45 50	
ctg agc ttc aac tat atc agg aca gtc act gct tca tcc ttc ccc ttt	907
Leu Ser Phe Asn Tyr Ile Arg Thr Val Thr Ala Ser Ser Phe Pro Phe	
55 60 65	
ctg gaa cag ctg cag ctg ctg gag ctc ggg agc cag tat acc ccc ttg	955
Leu Glu Gln Leu Gln Leu Leu Glu Leu Gly Ser Gln Tyr Thr Pro Leu	
70 75 80	
act att gac aag gag gcc ttc aga aac ctg ccc aac ctt aga atc ttg	1003
Thr Ile Asp Lys Glu Ala Phe Arg Asn Leu Pro Asn Leu Arg Ile Leu	
85 90 95 100	
gac ctg gga agt agt aag ata tac ttc ttg cat cca gat gct ttt cag	1051
Asp Leu Gly Ser Ser Lys Ile Tyr Phe Leu His Pro Asp Ala Phe Gln	
105 110 115	
gga ctg ttc cat ctg ttt gaa ctt aga ctg tat ttc tgt ggt ctc tct	1099
Gly Leu Phe His Leu Phe Glu Leu Arg Leu Tyr Phe Cys Gly Leu Ser	
120 125 130	
gat gct gta ttg aaa gat ggt tat ttc aga aat tta aag gct tta act	1147
Asp Ala Val Leu Lys Asp Gly Tyr Phe Arg Asn Leu Lys Ala Leu Thr	
135 140 145	
cgc ttg gat cta tcc aaa aat cag att cgt agc ctt tac ctt cat cct	1195
Arg Leu Asp Leu Ser Lys Asn Gln Ile Arg Ser Leu Tyr Leu His Pro	
150 155 160	
tca ttt ggg aag ttg aat tcc tta aag tcc ata gat ttt tcc tcc aac	1243
Ser Phe Gly Lys Leu Asn Ser Leu Lys Ser Ile Asp Phe Ser Ser Asn	
165 170 175 180	
caa ata ttc ctt gta tgt gaa cat gag ctc gag ccc cta caa ggg aaa	1291
Gln Ile Phe Leu Val Cys Glu His Glu Leu Glu Pro Leu Gln Gly Lys	
185 190 195	
acg ctc tcc ttt ttt agc ctc gca gct aat agc ttg tat agc aga gtc	1339
Thr Leu Ser Phe Phe Ser Leu Ala Ala Asn Ser Leu Tyr Ser Arg Val	
200 205 210	

tca gtg gac tgg gga aaa tgt atg aac cca ttc aga aac atg gtg ctg	1387
Ser Val Asp Trp Gly Lys Cys Met Asn Pro Phe Arg Asn Met Val Leu	
215 220 225	
gag ata gta gat gtt tct gga aat ggc tgg aca gtg gac atc aca gga	1435
Glu Ile Val Asp Val Ser Gly Asn Gly Trp Thr Val Asp Ile Thr Gly	
230 235 240	
aac ttt agc aat gcc atc agc aaa agc cag gcc ttc tct ttg att ctt	1483
Asn Phe Ser Asn Ala Ile Ser Lys Ser Gln Ala Phe Ser Leu Ile Leu	
245 250 255 260	
gcc cac cac atc atg ggt gcc ggg ttt ggc ttc cat aac atc aaa gat	1531
Ala His His Ile Met Gly Ala Gly Phe Gly Phe His Asn Ile Lys Asp	
265 270 275	
cct gac cag aac aca ttt gct ggc ctg gcc aga agt tca gtg aga cac	1579
Pro Asp Gln Asn Thr Phe Ala Gly Leu Ala Arg Ser Ser Val Arg His	
280 285 290	
ctg gac ctt tca cat ggg ttt gtc ttc tcc ctg aac tca cga gtc ttt	1627
Leu Asp Leu Ser His Gly Phe Val Phe Ser Leu Asn Ser Arg Val Phe	
295 300 305	
gag aca ctc aag gat ttg aag gtt ctg aac ctt gcc tac aac aag ata	1675
Glu Thr Leu Lys Asp Leu Lys Val Leu Asn Leu Ala Tyr Asn Lys Ile	
310 315 320	
aat aag att gca gat gaa gca ttt tac gga ctt gac aac ctc caa gtt	1723
Asn Lys Ile Ala Asp Glu Ala Phe Tyr Gly Leu Asp Asn Leu Gln Val	
325 330 335 340	
ctc aat ttg tca tat aac ctt ctg ggg gaa ctt tgc agt tcg aat ttc	1771
Leu Asn Leu Ser Tyr Asn Leu Leu Gly Glu Leu Cys Ser Ser Asn Phe	
345 350 355	
tat gga cta cct aag gta gcc tac att gat ttg caa aag aat cac att	1819
Tyr Gly Leu Pro Lys Val Ala Tyr Ile Asp Leu Gln Lys Asn His Ile	
360 365 370	
gca ata att caa gac caa aca ttc aaa ttc ctg gaa aaa tta cag acc	1867
Ala Ile Ile Gln Asp Gln Thr Phe Lys Phe Leu Glu Lys Leu Gln Thr	
375 380 385	
ttg gat ctc cga gac aat gct ctt aca acc att cat ttt att cca agc	1915
Leu Asp Leu Arg Asp Asn Ala Leu Thr Thr Ile His Phe Ile Pro Ser	
390 395 400	
ata ccc gat atc ttc ttg agt ggc aat aaa cta gtg act ttg cca aag	1963
Ile Pro Asp Ile Phe Leu Ser Gly Asn Lys Leu Val Thr Leu Pro Lys	
405 410 415 420	
atc aac ctt aca gcg aac ctc atc cac tta tca gaa aac agg cta gaa	2011
Ile Asn Leu Thr Ala Asn Leu Ile His Leu Ser Glu Asn Arg Leu Glu	

	425	430	435	
aat cta gat att ctc tac ttt ctc cta cgg gta cct cat ctc cag att				2059
Asn Leu Asp Ile Leu Tyr Phe Leu Leu Arg Val Pro His Leu Gln Ile				
	440	445	450	
ctc att tta aat caa aat cgc ttc tcc tcc tgt agt gga gat caa acc				2107
Leu Ile Leu Asn Gln Asn Arg Phe Ser Ser Cys Ser Gly Asp Gln Thr				
	455	460	465	
cct tca gag aat ccc agc tta gaa cag ctt ttc ctt gga gaa aat atg				2155
Pro Ser Glu Asn Pro Ser Leu Glu Gln Leu Phe Leu Gly Glu Asn Met				
	470	475	480	
ttg caa ctt gcc tgg gaa act gag ctc tgt tgg gat gtt ttt gag gga				2203
Leu Gln Leu Ala Trp Glu Thr Glu Leu Cys Trp Asp Val Phe Glu Gly				
	485	490	495	500
ctt tct cat ctt caa gtt ctg tat ttg aat cat aac tat ctt aat tcc				2251
Leu Ser His Leu Gln Val Leu Tyr Leu Asn His Asn Tyr Leu Asn Ser				
	505	510	515	
ctt cca cca gga gta ttt agc cat ctg act gca tta agg gga cta agc				2299
Leu Pro Pro Gly Val Phe Ser His Leu Thr Ala Leu Arg Gly Leu Ser				
	520	525	530	
ctc aac tcc aac agg ctg aca gtt ctt tct cac aat gat tta cct gct				2347
Leu Asn Ser Asn Arg Leu Thr Val Leu Ser His Asn Asp Leu Pro Ala				
	535	540	545	
aat tta gag atc ctg gac ata tcc agg aac cag ctc cta gct cct aat				2395
Asn Leu Glu Ile Leu Asp Ile Ser Arg Asn Gln Leu Leu Ala Pro Asn				
	550	555	560	
cct gat gta ttt gta tca ctt agt gtc ttg gat ata act cat aac aag				2443
Pro Asp Val Phe Val Ser Leu Ser Val Leu Asp Ile Thr His Asn Lys				
	565	570	575	580
ttc att tgt gaa tgt gaa ctt agc act ttt atc aat tgg ctt aat cac				2491
Phe Ile Cys Glu Cys Glu Leu Ser Thr Phe Ile Asn Trp Leu Asn His				
	585	590	595	
acc aat gtc act ata gct ggg cct cct gca gac ata tat tgt gtg tac				2539
Thr Asn Val Thr Ile Ala Gly Pro Pro Ala Asp Ile Tyr Cys Val Tyr				
	600	605	610	
cct gac tgg ctc tct ggg gtt tcc ctc ttc tct ctt tcc acg gaa ggt				2587
Pro Asp Ser Leu Ser Gly Val Ser Leu Phe Ser Leu Ser Thr Glu Gly				
	615	620	625	
tgt gat gaa gag gaa gtc tta aag tcc cta aag ttc tcc ctt ttc att				2635
Cys Asp Glu Glu Glu Val Leu Lys Ser Leu Lys Phe Ser Leu Phe Ile				
	630	635	640	
gta tgc act gtc act ctg act ctg ttc ctc atg acc atc ctc aca gtc				2683

Val Cys Thr Val Thr Leu Thr Leu Phe Leu Met Thr Ile Leu Thr Val	
645	650 655 660
aca aag ttc cgg ggc ttc tgt ttt atc tgt tat aag aca gcc cag aga	2731
Thr Lys Phe Arg Gly Phe Cys Phe Ile Cys Tyr Lys Thr Ala Gln Arg	
665	670 675
ctg gtg ttc aag gac cat ccc cag ggc aca gaa cct gat atg tac aaa	2779
Leu Val Phe Lys Asp His Pro Gln Gly Thr Glu Pro Asp Met Tyr Lys	
680	685 690
tat gat gcc tat ttg tgc ttc agc agc aaa gac ttc aca tgg gtg cag	2827
Tyr Asp Ala Tyr Leu Cys Phe Ser Ser Lys Asp Phe Thr Trp Val Gln	
695	700 705
aat gct ttg ctc aaa cac ctg gac act caa tac agt gac caa aac aga	2875
Asn Ala Leu Leu Lys His Leu Asp Thr Gln Tyr Ser Asp Gln Asn Arg	
710	715 720
ttc aac ctg tgc ttt gaa gaa aga gac ttt gtc cca gga gaa aac cgc	2923
Phe Asn Leu Cys Phe Glu Glu Arg Asp Phe Val Pro Gly Glu Asn Arg	
725	730 735 740
att gcc aat atc cag gat gcc atc tgg aac agt aga aag atc gtt tgt	2971
Ile Ala Asn Ile Gln Asp Ala Ile Trp Asn Ser Arg Lys Ile Val Cys	
745	750 755
ctt gtg agc aga cac ttc ctt aga gat ggc tgg tgc ctt gaa gcc ttc	3019
Leu Val Ser Arg His Phe Leu Arg Asp Gly Trp Cys Leu Glu Ala Phe	
760	765 770
agt tat gcc cag ggc agg tgc tta tct gac ctt aac agt gct ctc atc	3067
Ser Tyr Ala Gln Gly Arg Cys Leu Ser Asp Leu Asn Ser Ala Leu Ile	
775	780 785
atg gtg gtg gtt ggg tcc ttg tcc cag tac cag ttg atg aaa cat caa	3115
Met Val Val Val Gly Ser Leu Ser Gln Tyr Gln Leu Met Lys His Gln	
790	795 800
tcc atc aga ggc ttt gta cag aaa cag cag tat ttg agg tgg cct gag	3163
Ser Ile Arg Gly Phe Val Gln Lys Gln Gln Tyr Leu Arg Trp Pro Glu	
805	810 815 820
gat ctc cag gat gtt ggc tgg ttt ctt cat aaa ctc tct caa cag ata	3211
Asp Leu Gln Asp Val Gly Trp Phe Leu His Lys Leu Ser Gln Gln Ile	
825	830 835
cta aag aaa gaa aaa gaa aag aag aaa gac aat aac att ccg ttg caa	3259
Leu Lys Lys Glu Lys Glu Lys Lys Lys Asp Asn Asn Ile Pro Leu Gln	
840	845 850
act gta gca acc atc tcc taatcaaagg agcaatttcc aacttatctc	3307
Thr Val Ala Thr Ile Ser	
855	

aagccacaaa taactcttca ctttgatatt gcaccaagtt atcatttttg ggtcctctct 3367
 ggaggttttt tttttctttt tgctactatg aaaacaacat aaatctctca attttcgat 3427
 caaa 3431

<210> 6

<211> 858

<212> PRT

<213> Homo sapiens

<400> 6

Met	Gly	Asp	His	Leu	Asp	Leu	Leu	Leu	Gly	Val	Val	Leu	Met	Ala	Gly
1				5					10					15	
Pro	Val	Phe	Gly	Ile	Pro	Ser	Cys	Ser	Phe	Asp	Gly	Arg	Ile	Ala	Phe
		20					25					30			
Tyr	Arg	Phe	Cys	Asn	Leu	Thr	Gln	Val	Pro	Gln	Val	Leu	Asn	Thr	Thr
	35					40					45				
Glu	Arg	Leu	Leu	Leu	Ser	Phe	Asn	Tyr	Ile	Arg	Thr	Val	Thr	Ala	Ser
	50					55					60				
Ser	Phe	Pro	Phe	Leu	Glu	Gln	Leu	Gln	Leu	Leu	Glu	Leu	Gly	Ser	Gln
65				70					75					80	
Tyr	Thr	Pro	Leu	Thr	Ile	Asp	Lys	Glu	Ala	Phe	Arg	Asn	Leu	Pro	Asn
			85					90					95		
Leu	Arg	Ile	Leu	Asp	Leu	Gly	Ser	Ser	Lys	Ile	Tyr	Phe	Leu	His	Pro
			100				105						110		
Asp	Ala	Phe	Gln	Gly	Leu	Phe	His	Leu	Phe	Glu	Leu	Arg	Leu	Tyr	Phe
	115				120						125				
Cys	Gly	Leu	Ser	Asp	Ala	Val	Leu	Lys	Asp	Gly	Tyr	Phe	Arg	Asn	Leu
	130				135						140				
Lys	Ala	Leu	Thr	Arg	Leu	Asp	Leu	Ser	Lys	Asn	Gln	Ile	Arg	Ser	Leu
145				150					155					160	
Tyr	Leu	His	Pro	Ser	Phe	Gly	Lys	Leu	Asn	Ser	Leu	Lys	Ser	Ile	Asp
			165					170						175	
Phe	Ser	Ser	Asn	Gln	Ile	Phe	Leu	Val	Cys	Glu	His	Glu	Leu	Glu	Pro
			180				185					190			
Leu	Gln	Gly	Lys	Thr	Leu	Ser	Phe	Phe	Ser	Leu	Ala	Ala	Asn	Ser	Leu
	195					200						205			
Tyr	Ser	Arg	Val	Ser	Val	Asp	Trp	Gly	Lys	Cys	Met	Asn	Pro	Phe	Arg
	210				215						220				
Asn	Met	Val	Leu	Glu	Ile	Val	Asp	Val	Ser	Gly	Asn	Gly	Trp	Thr	Val
225				230						235				240	
Asp	Ile	Thr	Gly	Asn	Phe	Ser	Asn	Ala	Ile	Ser	Lys	Ser	Gln	Ala	Phe
			245					250					255		
Ser	Leu	Ile	Leu	Ala	His	His	Ile	Met	Gly	Ala	Gly	Phe	Gly	Phe	His
		260					265					270			
Asn	Ile	Lys	Asp	Pro	Asp	Gln	Asn	Thr	Phe	Ala	Gly	Leu	Ala	Arg	Ser
	275					280						285			
Ser	Val	Arg	His	Leu	Asp	Leu	Ser	His	Gly	Phe	Val	Phe	Ser	Leu	Asn
	290					295					300				
Ser	Arg	Val	Phe	Glu	Thr	Leu	Lys	Asp	Leu	Lys	Val	Leu	Asn	Leu	Ala
305				310						315				320	
Tyr	Asn	Lys	Ile	Asn	Lys	Ile	Ala	Asp	Glu	Ala	Phe	Tyr	Gly	Leu	Asp
			325					330					335		
Asn	Leu	Gln	Val	Leu	Asn	Leu	Ser	Tyr	Asn	Leu	Leu	Gly	Glu	Leu	Cys
	340					345						350			
Ser	Ser	Asn	Phe	Tyr	Gly	Leu	Pro	Lys	Val	Ala	Tyr	Ile	Asp	Leu	Gln

355	360	365
Lys Asn His Ile Ala Ile Ile Gln Asp Gln Thr Phe Lys Phe Leu Glu		
370	375	380
Lys Leu Gln Thr Leu Asp Leu Arg Asp Asn Ala Leu Thr Thr Ile His		
385	390	395
Phe Ile Pro Ser Ile Pro Asp Ile Phe Leu Ser Gly Asn Lys Leu Val		
405	410	415
Thr Leu Pro Lys Ile Asn Leu Thr Ala Asn Leu Ile His Leu Ser Glu		
420	425	430
Asn Arg Leu Glu Asn Leu Asp Ile Leu Tyr Phe Leu Leu Arg Val Pro		
435	440	445
His Leu Gln Ile Leu Ile Leu Asn Gln Asn Arg Phe Ser Ser Cys Ser		
450	455	460
Gly Asp Gln Thr Pro Ser Glu Asn Pro Ser Leu Glu Gln Leu Phe Leu		
465	470	475
Gly Glu Asn Met Leu Gln Leu Ala Trp Glu Thr Glu Leu Cys Trp Asp		
485	490	495
Val Phe Glu Gly Leu Ser His Leu Gln Val Leu Tyr Leu Asn His Asn		
500	505	510
Tyr Leu Asn Ser Leu Pro Pro Gly Val Phe Ser His Leu Thr Ala Leu		
515	520	525
Arg Gly Leu Ser Leu Asn Ser Asn Arg Leu Thr Val Leu Ser His Asn		
530	535	540
Asp Leu Pro Ala Asn Leu Glu Ile Leu Asp Ile Ser Arg Asn Gln Leu		
545	550	555
Leu Ala Pro Asn Pro Asp Val Phe Val Ser Leu Ser Val Leu Asp Ile		
565	570	575
Thr His Asn Lys Phe Ile Cys Glu Cys Glu Leu Ser Thr Phe Ile Asn		
580	585	590
Trp Leu Asn His Thr Asn Val Thr Ile Ala Gly Pro Pro Ala Asp Ile		
595	600	605
Tyr Cys Val Tyr Pro Asp Ser Leu Ser Gly Val Ser Leu Phe Ser Leu		
610	615	620
Ser Thr Glu Gly Cys Asp Glu Glu Glu Val Leu Lys Ser Leu Lys Phe		
625	630	635
Ser Leu Phe Ile Val Cys Thr Val Thr Leu Thr Leu Phe Leu Met Thr		
645	650	655
Ile Leu Thr Val Thr Lys Phe Arg Gly Phe Cys Phe Ile Cys Tyr Lys		
660	665	670
Thr Ala Gln Arg Leu Val Phe Lys Asp His Pro Gln Gly Thr Glu Pro		
675	680	685
Asp Met Tyr Lys Tyr Asp Ala Tyr Leu Cys Phe Ser Ser Lys Asp Phe		
690	695	700
Thr Trp Val Gln Asn Ala Leu Leu Lys His Leu Asp Thr Gln Tyr Ser		
705	710	715
Asp Gln Asn Arg Phe Asn Leu Cys Phe Glu Glu Arg Asp Phe Val Pro		
725	730	735
Gly Glu Asn Arg Ile Ala Asn Ile Gln Asp Ala Ile Trp Asn Ser Arg		
740	745	750
Lys Ile Val Cys Leu Val Ser Arg His Phe Leu Arg Asp Gly Trp Cys		
755	760	765
Leu Glu Ala Phe Ser Tyr Ala Gln Gly Arg Cys Leu Ser Asp Leu Asn		
770	775	780
Ser Ala Leu Ile Met Val Val Val Gly Ser Leu Ser Gln Tyr Gln Leu		
785	790	795
		800

Met Lys His Gln Ser Ile Arg Gly Phe Val Gln Lys Gln Gln Tyr Leu
 805 810 815
 Arg Trp Pro Glu Asp Leu Gln Asp Val Gly Trp Phe Leu His Lys Leu
 820 825 830
 Ser Gln Gln Ile Leu Lys Lys Glu Lys Glu Lys Lys Lys Asp Asn Asn
 835 840 845
 Ile Pro Leu Gln Thr Val Ala Thr Ile Ser
 850 855

<210> 7

<211> 1839

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)...(1836)

<223> Chimera Mus musculus and Homo sapiens

<400> 7

atg tgc cga gcc atc tct ctt agg cgc ttg ctg ctg ctg ctg ctg cag 48
 Met Cys Arg Ala Ile Ser Leu Arg Arg Leu Leu Leu Leu Leu Leu Gln
 1 5 10 15

ctg tca caa ctc cta gct gtc act caa ggg aag acg ctg gtg ctg ggg 96
 Leu Ser Gln Leu Leu Ala Val Thr Gln Gly Lys Thr Leu Val Leu Gly
 20 25 30

aag gaa ggg gaa tca gca gaa ctg ccc tgc gag agt tcc cag aag aag 144
 Lys Glu Gly Glu Ser Ala Glu Leu Pro Cys Glu Ser Ser Gln Lys Lys
 35 40 45

atc aca gtc ttc acc tgg aag ttc tct gac cag agg aag att ctg ggg 192
 Ile Thr Val Phe Thr Trp Lys Phe Ser Asp Gln Arg Lys Ile Leu Gly
 50 55 60

cag cat ggc aaa ggt gta tta att aga gga ggt tgc cct tgc cag ttt 240
 Gln His Gly Lys Gly Val Leu Ile Arg Gly Gly Ser Pro Ser Gln Phe
 65 70 75 80

gat cgt ttt gat tcc aaa aaa ggg gca tgg gag aaa gga tgc ttt cct 288
 Asp Arg Phe Asp Ser Lys Lys Gly Ala Trp Glu Lys Gly Ser Phe Pro
 85 90 95

ctc atc atc aat aaa ctt aag atg gaa gac tct cag act tat atc tgt 336
 Leu Ile Ile Asn Lys Leu Lys Met Glu Asp Ser Gln Thr Tyr Ile Cys
 100 105 110

gag ctg gag aac agg aaa gag gag gtg gag ttg tgg gtg ttc aaa gtg 384
 Glu Leu Glu Asn Arg Lys Glu Glu Val Glu Leu Trp Val Phe Lys Val
 115 120 125

acc ttc agt ccg ggt acc agc ctg ttg caa ggg cag agc ctg acc ctg 432

Thr	Phe	Ser	Pro	Gly	Thr	Ser	Leu	Leu	Gln	Gly	Gln	Ser	Leu	Thr	Leu		
130						135					140						
acc	ttg	gat	agc	aac	tct	aag	gtc	tct	aac	ccc	ttg	aca	gag	tgc	aaa	480	
Thr	Leu	Asp	Ser	Asn	Ser	Lys	Val	Ser	Asn	Pro	Leu	Thr	Glu	Cys	Lys		
145					150				155						160		
cac	aaa	aag	ggc	aaa	gtt	gtc	agt	ggc	tcc	aaa	gtt	ctc	tcc	atg	tcc	528	
His	Lys	Lys	Gly	Lys	Val	Val	Ser	Gly	Ser	Lys	Val	Leu	Ser	Met	Ser		
				165				170						175			
aac	cta	agg	gtt	cag	gac	agc	gac	ttc	tgg	aac	tgc	acc	gtg	acc	ctg	576	
Asn	Leu	Arg	Val	Gln	Asp	Ser	Asp	Phe	Trp	Asn	Cys	Thr	Val	Thr	Leu		
			180					185					190				
gac	cag	aaa	aag	aac	tgg	ttc	ggc	atg	aca	ctc	tca	gtg	ctg	ggc	ttt	624	
Asp	Gln	Lys	Lys	Asn	Trp	Phe	Gly	Met	Thr	Leu	Ser	Val	Leu	Gly	Phe		
		195				200						205					
cag	agc	aca	gct	atc	acg	gcc	tat	aag	agt	gag	gga	gag	tca	gcg	gag	672	
Gln	Ser	Thr	Ala	Ile	Thr	Ala	Tyr	Lys	Ser	Glu	Gly	Glu	Ser	Ala	Glu		
		210				215					220						
ttc	tcc	ttc	cca	ctc	aac	ttt	gca	gag	gaa	aac	ggg	tgg	gga	gag	ctg	720	
Phe	Ser	Phe	Pro	Leu	Asn	Phe	Ala	Glu	Glu	Asn	Gly	Trp	Gly	Glu	Leu		
225					230					235					240		
atg	tgg	aag	gca	gag	aag	gat	tct	ttc	ttc	cag	ccc	tgg	atc	tcc	ttc	768	
Met	Trp	Lys	Ala	Glu	Lys	Asp	Ser	Phe	Phe	Gln	Pro	Trp	Ile	Ser	Phe		
				245				250						255			
tcc	ata	aag	aac	aaa	gag	gtg	tcc	gta	caa	aag	tcc	acc	aaa	gac	ctc	816	
Ser	Ile	Lys	Asn	Lys	Glu	Val	Ser	Val	Gln	Lys	Ser	Thr	Lys	Asp	Leu		
			260					265					270				
aag	ctc	cag	ctg	aag	gaa	acg	ctc	cca	ctc	acc	ctc	aag	ata	ccc	cag	864	
Lys	Leu	Gln	Leu	Lys	Glu	Thr	Leu	Pro	Leu	Thr	Leu	Lys	Ile	Pro	Gln		
			275				280					285					
gtc	tgc	ctt	cag	ttt	gct	ggc	tct	ggc	aac	ctg	act	ctg	act	ctg	gac	912	
Val	Ser	Leu	Gln	Phe	Ala	Gly	Ser	Gly	Asn	Leu	Thr	Leu	Thr	Leu	Asp		
		290				295					300						
aaa	ggg	aca	ctg	cat	cag	gaa	gtg	aac	ctg	gtg	gtg	atg	aaa	gtg	gct	960	
Lys	Gly	Thr	Leu	His	Gln	Glu	Val	Asn	Leu	Val	Val	Met	Lys	Val	Ala		
305					310				315						320		
cag	ctc	aac	aat	act	ttg	acc	tgt	gag	gtg	atg	gga	cct	acc	tct	ccc	1008	
Gln	Leu	Asn	Asn	Thr	Leu	Thr	Cys	Glu	Val	Met	Gly	Pro	Thr	Ser	Pro		
				325					330					335			
aag	atg	aga	ctg	acc	ctg	aag	cag	gag	aac	cag	gag	gcc	agg	gtc	tct	1056	
Lys	Met	Arg	Leu	Thr	Leu	Lys	Gln	Glu	Asn	Gln	Glu	Ala	Arg	Val	Ser		
			340					345					350				

gag gag cag aaa gta gtt caa gtg gtg gcc cct gag aca ggg ctg tgg	1104
Glu Glu Gln Lys Val Val Gln Val Val Ala Pro Glu Thr Gly Leu Trp	
355 360 365	
cag tgt cta ctg agt gaa ggt gat aag gtc aag atg gac tcc agg atc	1152
Gln Cys Leu Leu Ser Glu Gly Asp Lys Val Lys Met Asp Ser Arg Ile	
370 375 380	
cag gtt tta tcc aga ggg gtg tac cag ttc tcc ctt ttc att gta tgc	1200
Gln Val Leu Ser Arg Gly Val Tyr Gln Phe Ser Leu Phe Ile Val Cys	
385 390 395 400	
act gtc act ctg act ctg ttc ctc atg acc atc ctc aca gtc aca aag	1248
Thr Val Thr Leu Thr Leu Phe Leu Met Thr Ile Leu Thr Val Thr Lys	
405 410 415	
ttc cgg ggc ttc tgt ttt atc tgt tat aag aca gcc cag aga ctg gtg	1296
Phe Arg Gly Phe Cys Phe Ile Cys Tyr Lys Thr Ala Gln Arg Leu Val	
420 425 430	
ttc aag gac cat ccc cag ggc aca gaa cct gat atg tac aaa tat gat	1344
Phe Lys Asp His Pro Gln Gly Thr Glu Pro Asp Met Tyr Lys Tyr Asp	
435 440 445	
gcc tat ttg tgc ttc agc agc aaa gac ttc aca tgg gtg cag aat gct	1392
Ala Tyr Leu Cys Phe Ser Ser Lys Asp Phe Thr Trp Val Gln Asn Ala	
450 455 460	
ttg ctc aaa cac ctg gac act caa tac agt gac caa aac aga ttc aac	1440
Leu Leu Lys His Leu Asp Thr Gln Tyr Ser Asp Gln Asn Arg Phe Asn	
465 470 475 480	
ctg tgc ttt gaa gaa aga gac ttt gtc cca gga gaa aac cgc att gcc	1488
Leu Cys Phe Glu Glu Arg Asp Phe Val Pro Gly Glu Asn Arg Ile Ala	
485 490 495	
aat atc cag gat gcc atc tgg aac agt aga aag atc gtt tgt ctt gtg	1536
Asn Ile Gln Asp Ala Ile Trp Asn Ser Arg Lys Ile Val Cys Leu Val	
500 505 510	
agc aga cac ttc ctt aga gat ggc tgg tgc ctt gaa gcc ttc agt tat	1584
Ser Arg His Phe Leu Arg Asp Gly Trp Cys Leu Glu Ala Phe Ser Tyr	
515 520 525	
gcc cag ggc agg tgc tta tct gac ctt aac agt gct ctc atc atg gtg	1632
Ala Gln Gly Arg Cys Leu Ser Asp Leu Asn Ser Ala Leu Ile Met Val	
530 535 540	
gtg gtt ggg tcc ttg tcc cag tac cag ttg atg aaa cat caa tcc atc	1680
Val Val Gly Ser Leu Ser Gln Tyr Gln Leu Met Lys His Gln Ser Ile	
545 550 555 560	
aga ggc ttt gta cag aaa cag cag tat ttg agg tgg cct gag gat ctc	1728
Arg Gly Phe Val Gln Lys Gln Gln Tyr Leu Arg Trp Pro Glu Asp Leu	
565 570 575	

cag gat gtt ggc tgg ttt ctt cat aaa ctc tct caa cag ata cta aag 1776
 Gln Asp Val Gly Trp Phe Leu His Lys Leu Ser Gln Gln Ile Leu Lys
 580 585 590

aaa gaa aaa gaa aag aag aaa gac aat aac att ccg ttg caa act gta 1824
 Lys Glu Lys Glu Lys Lys Lys Asp Asn Asn Ile Pro Leu Gln Thr Val
 595 600 605

gca acc atc tcc taa 1839
 Ala Thr Ile Ser
 610

<210> 8
 <211> 612
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Chimera Mus musculus and Homo sapiens

<400> 8
 Met Cys Arg Ala Ile Ser Leu Arg Arg Leu Leu Leu Leu Leu Leu Gln
 1 5 10 15
 Leu Ser Gln Leu Leu Ala Val Thr Gln Gly Lys Thr Leu Val Leu Gly
 20 25 30
 Lys Glu Gly Glu Ser Ala Glu Leu Pro Cys Glu Ser Ser Gln Lys Lys
 35 40 45
 Ile Thr Val Phe Thr Trp Lys Phe Ser Asp Gln Arg Lys Ile Leu Gly
 50 55 60
 Gln His Gly Lys Gly Val Leu Ile Arg Gly Gly Ser Pro Ser Gln Phe
 65 70 75 80
 Asp Arg Phe Asp Ser Lys Lys Gly Ala Trp Glu Lys Gly Ser Phe Pro
 85 90 95
 Leu Ile Ile Asn Lys Leu Lys Met Glu Asp Ser Gln Thr Tyr Ile Cys
 100 105 110
 Glu Leu Glu Asn Arg Lys Glu Glu Val Glu Leu Trp Val Phe Lys Val
 115 120 125
 Thr Phe Ser Pro Gly Thr Ser Leu Leu Gln Gly Gln Ser Leu Thr Leu
 130 135 140
 Thr Leu Asp Ser Asn Ser Lys Val Ser Asn Pro Leu Thr Glu Cys Lys
 145 150 155 160
 His Lys Lys Gly Lys Val Val Ser Gly Ser Lys Val Leu Ser Met Ser
 165 170 175
 Asn Leu Arg Val Gln Asp Ser Asp Phe Trp Asn Cys Thr Val Thr Leu
 180 185 190
 Asp Gln Lys Lys Asn Trp Phe Gly Met Thr Leu Ser Val Leu Gly Phe
 195 200 205
 Gln Ser Thr Ala Ile Thr Ala Tyr Lys Ser Glu Gly Glu Ser Ala Glu
 210 215 220
 Phe Ser Phe Pro Leu Asn Phe Ala Glu Glu Asn Gly Trp Gly Glu Leu
 225 230 235 240
 Met Trp Lys Ala Glu Lys Asp Ser Phe Phe Gln Pro Trp Ile Ser Phe
 245 250 255

Ser Ile Lys Asn Lys Glu Val Ser Val Gln Lys Ser Thr Lys Asp Leu
 260 265 270
 Lys Leu Gln Leu Lys Glu Thr Leu Pro Leu Thr Leu Lys Ile Pro Gln
 275 280 285
 Val Ser Leu Gln Phe Ala Gly Ser Gly Asn Leu Thr Leu Thr Leu Asp
 290 295 300
 Lys Gly Thr Leu His Gln Glu Val Asn Leu Val Val Met Lys Val Ala
 305 310 315 320
 Gln Leu Asn Asn Thr Leu Thr Cys Glu Val Met Gly Pro Thr Ser Pro
 325 330 335
 Lys Met Arg Leu Thr Leu Lys Gln Glu Asn Gln Glu Ala Arg Val Ser
 340 345 350
 Glu Glu Gln Lys Val Val Gln Val Val Ala Pro Glu Thr Gly Leu Trp
 355 360 365
 Gln Cys Leu Leu Ser Glu Gly Asp Lys Val Lys Met Asp Ser Arg Ile
 370 375 380
 Gln Val Leu Ser Arg Gly Val Tyr Gln Phe Ser Leu Phe Ile Val Cys
 385 390 395 400
 Thr Val Thr Leu Thr Leu Phe Leu Met Thr Ile Leu Thr Val Thr Lys
 405 410 415
 Phe Arg Gly Phe Cys Phe Ile Cys Tyr Lys Thr Ala Gln Arg Leu Val
 420 425 430
 Phe Lys Asp His Pro Gln Gly Thr Glu Pro Asp Met Tyr Lys Tyr Asp
 435 440 445
 Ala Tyr Leu Cys Phe Ser Ser Lys Asp Phe Thr Trp Val Gln Asn Ala
 450 455 460
 Leu Leu Lys His Leu Asp Thr Gln Tyr Ser Asp Gln Asn Arg Phe Asn
 465 470 475 480
 Leu Cys Phe Glu Glu Arg Asp Phe Val Pro Gly Glu Asn Arg Ile Ala
 485 490 495
 Asn Ile Gln Asp Ala Ile Trp Asn Ser Arg Lys Ile Val Cys Leu Val
 500 505 510
 Ser Arg His Phe Leu Arg Asp Gly Trp Cys Leu Glu Ala Phe Ser Tyr
 515 520 525
 Ala Gln Gly Arg Cys Leu Ser Asp Leu Asn Ser Ala Leu Ile Met Val
 530 535 540
 Val Val Gly Ser Leu Ser Gln Tyr Gln Leu Met Lys His Gln Ser Ile
 545 550 555 560
 Arg Gly Phe Val Gln Lys Gln Gln Tyr Leu Arg Trp Pro Glu Asp Leu
 565 570 575
 Gln Asp Val Gly Trp Phe Leu His Lys Leu Ser Gln Gln Ile Leu Lys
 580 585 590
 Lys Glu Lys Glu Lys Lys Lys Asp Asn Asn Ile Pro Leu Gln Thr Val
 595 600 605
 Ala Thr Ile Ser
 610

INTERNATIONAL SEARCH REPORT

International Application No
PCT/US 01/22978

A. CLASSIFICATION OF SUBJECT MATTER
IPC 7 C07K14/255 C12N15/31 A61K39/00

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 7 C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the International search (name of data base and, where practical, search terms used)

BIOSIS, MEDLINE, EPO-Internal, WPI Data, PAJ, CHEM ABS Data

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>NEWTON S M C ET AL: "IMMUNE RESPONSE TO CHOLERA TOXIN EPITOPE INSERTED IN SALMONELLA FLAGELLIN" SCIENCE (WASHINGTON D C), vol. 244, no. 4900, 1989, pages 70-72, XP001094029 ISSN: 0036-8075 the whole document</p> <p style="text-align: center;">--- -/--</p>	1-6, 11-22

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

* Special categories of cited documents:

- *A* document defining the general state of the art which is not considered to be of particular relevance
- *E* earlier document but published on or after the international filing date
- *L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- *O* document referring to an oral disclosure, use, exhibition or other means
- *P* document published prior to the international filing date but later than the priority date claimed

- *T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- *X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- *Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- *A* document member of the same patent family

Date of the actual completion of the international search

23 August 2002

Date of mailing of the international search report

11/09/2002

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax (+31-70) 340-3016

Authorized officer

Billang, J

INTERNATIONAL SEARCH REPORT

International Application No

PCT/US 01/22978

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	MCSORLEY S J ET AL: "Characterization of CD4+ T cell responses during natural infection with Salmonella typhimurium." JOURNAL OF IMMUNOLOGY (BALTIMORE, MD.: 1950) UNITED STATES 15 JAN 2000, vol. 164, no. 2, 15 January 2000 (2000-01-15), pages 986-993, XP002210840 ISSN: 0022-1767 the whole document ---	1-6,11, 12,15, 16,19-22
X	EAVES-PYLES T ET AL: "Flagellin, a novel mediator of Salmonella-induced epithelial activation and systemic inflammation: I kappa B alpha degradation, induction of nitric oxide synthase, induction of proinflammatory mediators, and cardiovascular dysfunction." JOURNAL OF IMMUNOLOGY (BALTIMORE, MD.: 1950) UNITED STATES 15 JAN 2001, vol. 166, no. 2, 15 January 2001 (2001-01-15), pages 1248-1260, XP002210841 ISSN: 0022-1767 the whole document ---	1-6,11, 12,15, 16,19-22
X	GEWIRTZ ANDREW T ET AL: "Salmonella typhimurium translocates flagellin across intestinal epithelia, inducing a proinflammatory response." JOURNAL OF CLINICAL INVESTIGATION, vol. 107, no. 1, January 2001 (2001-01), pages 99-109, XP002210842 ISSN: 0021-9738 the whole document ---	1-6,11, 12,15, 16,19-22
A	FELIX GEORG ET AL: "Plants have a sensitive perception system for the most conserved domain of bacterial flagellin" PLANT JOURNAL, BLACKWELL SCIENTIFIC PUBLICATIONS, OXFORD, GB, vol. 18, no. 3, May 1999 (1999-05), pages 265-276, XP002148873 ISSN: 0960-7412 ---	
P,X	HAYASHI FUMITAKA ET AL: "The innate immune response to bacterial flagellin is mediated by Toll-like receptor 5." NATURE (LONDON), vol. 410, no. 6832, 2001, pages 1099-1103, XP002210843 ISSN: 0028-0836 the whole document --- -/--	1-35

INTERNATIONAL SEARCH REPORT

ational Application No

PCT/US 01/22978

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
E	WO 02 09748 A (MEDZHITOV RUSLAN M PH D ;UNIV YALE (US)) 7 February 2002 (2002-02-07) page 51, line 21 -page 52, line 11 -----	5,6, 11-22

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 01/22978

Patent document cited in search report		Publication date		Patent family member(s)	Publication date
WO 0209748	A	07-02-2002	US	2002061312 A1	23-05-2002
			AU	8640501 A	13-02-2002
			WO	0209748 A1	07-02-2002